# BEST AVAILABLE COPY

(12) INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

(19) World Intellectual Property Organization
International Bureau





(43) International Publication Date 12 September 2003 (12.09.2003)

**PCT** 

# (10) International Publication Number WO 03/074654 A2

(51)	International Pate	ent Classification <sup>7</sup> :	C	l2N	US	60/363,124 (CON)
(21)	International App	lication Number:	PCT/US03/0	5028	Filed on US	11 March 2002 (11.03.2002) 60/386,782 (CON)
					Filed on US	6 June 2002 (06.06.2002) 60/406.784 (CON)
(22)	22) International Filing Date: 20 February 2003 (20.02.2003)				Filed on	29 August 2002 (29.08.2002)
(25)	Filing Language:		En	glish	US Filed on	60/408,378 (CON) 5 September 2002 (05.09.2002)
(26)	Publication Language:		En	glish	US Filed on	60/409,293 (CON) 9 September 2002 (09.09.2002)
(30)	Priority Data: 60/358,580	20 February 2002 (	(20.02.2002)	US	US Filed on	60/440,129 (CON) 15 January 2003 (15.01.2003)
	60/363,124 60/386,782 60/406,784	11 March 2002 ( 6 June 2002 ( 29 August 2002 (	(06.06.2002) (29.08.2002)	US US US		for all designated States except US): Sirna s, Inc [US/US]; 2950 Wilderness Place, 80301 (US).
	60/408,378	5 September 2002 (	(05.09.2002)	US		

US

US

(63) Related by continuation (CON) or continuation-in-part (CIP) to earlier applications:

US 60/358,580 (CON) Filed on 20 February 2002 (20.02.2002)

9 September 2002 (09.09.2002)

15 January 2003 (15.01.2003)

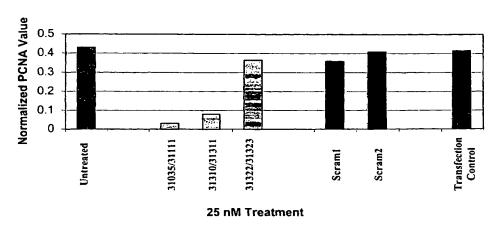
(72) Inventors; and

(75) Inventors/Applicants (for US only): MCSWIGGEN,
James [US/US]; 4866 Franklin Drive, Boulder, CO
80301 (US). BEIGELMAN, Leonid [US/US]; 5530
Colt Drive, Longmont, CO 80503 (US). CHOWRIRA,
Bharat [US/US]; 576 Manorwood Lane, Louisville, CO
80027 (US). PAVCO, Pamela [US/US]; 705 Barberry

[Continued on next page]

(54) Title: RNA INTERFERENCE MEDIATED INHIBITION OF GENE EXPRESSION USING SHORT INTERFERING NUCLEIC ACID (SINA)

## A549 24h PCNA mRNA Expression



(57) Abstract: The present invention concerns methods and reagents useful in modulating gene expression in a variety of applications, including use in therapeutic, diagnostic, target validation, and genomic discovery applications. Specifically, the invention relates to small nucleic acid molecules, such as short interfering nucleic acid (siNA), short interfering RNA (siRNA), double-stranded RNA (dsRNA), micro-RNA (miRNA), and short hairpin RNA (shRNA) molecules capable of mediating RNA interference (RNAi) against target nucleic acid sequences. The small nucleic acid molecules are useful in the treatment of any disease or condition that responds to modulation of gene expression or activity in a cell, tissue, or organism.

60/409,293

60/440,129



O 03/074654



Circle, Lafayette, CO 80026 (US). FOSNAUGH, Kathy [US/US]; 1030 Edinboro Drive, Boulder, Colorado 80305 (US). JAMISON, Sharon [US/US]; 4985 Twin Lakes Rd, #89, Boulder, CO 80301 (US). USMAN, Nassim [US/US]; 2129 Night Sky Lane, Lafayette, CO 80026 (US). THOMPSON, James [US/US]; 705 Barberry Circle, Lafayette, CO 80026 (US).

- (74) Agent: TERPSTRA, Anita, J.; McDonnell Boehnen Hulbert & Berghoff, 300 South Wacker Drive, Suite 3200, Chicago, IL 60606 (US).
- (81) Designated States (national): AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CO, CR, CU, CZ, DE, DK, DM, DZ, EC, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, OM, PH, PL, PT, RO, RU, SC, SD, SE, SG, SK, SL, TJ, TM, TN, TR, TT, TZ, UA, UG, US, UZ, VC, VN, YU, ZA, ZM, ZW.

(84) Designated States (regional): ARIPO patent (GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZM, ZW), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, BG, CH, CY, CZ, DE, DK, EE, ES, FI, FR, GB, GR, HU, IE, IT, LU, MC, NL, PT, SE, SI, SK, TR), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, GO, GW, ML, MR, NE, SN, TD, TG).

#### Declaration under Rule 4.17:

— of inventorship (Rule 4.17(iv)) for US only

#### Published:

 without international search report and to be republished upon receipt of that report

For two-letter codes and other abbreviations, refer to the "Guidance Notes on Codes and Abbreviations" appearing at the beginning of each regular issue of the PCT Gazette.

# RNA INTERFERENCE MEDIATED INHIBITION OF GENE EXPRESSION USING SHORT INTERFERING NUCLEIC ACID (siNA)

This invention claims the benefit of Beigelman USSN 60/358,580 filed February 20, 2002, of Beigelman USSN 60/363,124 filed March 11, 2002, of Beigelman USSN 60/386,782 filed June 6, 2002, of Beigelman USSN 60/406,784 filed August 29, 2002, of Beigelman USSN 60/408,378 filed September 5, 2002, of Beigelman USSN 60/409,293 filed September 9, 2002, and of Beigelman USSN 60/440,129 filed January 15, 2003. These applications are hereby incorporated by reference herein in their entireties, including the drawings.

10 Field Of The Invention

The present invention concerns methods and reagents useful in modulating gene expression in a variety of applications, including use in therapeutic, diagnostic, target validation, and genomic discovery applications. Specifically, the invention relates to small nucleic acid molecules, such as short interfering nucleic acid (siNA), short interfering RNA (siRNA), double-stranded RNA (dsRNA), micro-RNA (miRNA), and short hairpin RNA (shRNA) molecules capable of mediating RNA interference (RNAi).

### **Background Of The Invention**

The following is a discussion of relevant art pertaining to RNAi. The discussion is provided only for understanding of the invention that follows. The summary is not an admission that any of the work described below is prior art to the claimed invention. Applicant demonstrates herein that chemically modified short interfering nucleic acids possess the same capacity to mediate RNAi as do siRNA molecules and are expected to possess improved stability and activity in vivo; therefore, this discussion is not meant to be limiting only to siRNA and can be applied to siNA as a whole.

RNA interference refers to the process of sequence-specific post-transcriptional gene silencing in animals mediated by short interfering RNAs (siRNAs) (Fire *et al.*, 1998, *Nature*, 391, 806). The corresponding process in plants is commonly referred to as post-transcriptional gene silencing or RNA silencing and is also referred to as quelling in fungi. The process of post-transcriptional gene silencing is thought to be an

5

15

20

evolutionarily-conserved cellular defense mechanism used to prevent the expression of foreign genes and is commonly shared by diverse flora and phyla (Fire et al., 1999, Trends Genet., 15, 358). Such protection from foreign gene expression may have evolved in response to the production of double-stranded RNAs (dsRNAs) derived from viral infection or from the random integration of transposon elements into a host genome via a cellular response that specifically destroys homologous single-stranded RNA or viral genomic RNA. The presence of dsRNA in cells triggers the RNAi response though a mechanism that has yet to be fully characterized. This mechanism appears to be different from the interferon response that results from dsRNA-mediated activation of protein kinase PKR and 2',5'-oligoadenylate synthetase resulting in non-specific cleavage of mRNA by ribonuclease L.

The presence of long dsRNAs in cells stimulates the activity of a ribonuclease III enzyme referred to as dicer. Dicer is involved in the processing of the dsRNA into short pieces of dsRNA known as short interfering RNAs (siRNAs) (Berstein et al., 2001, Nature, 409, 363). Short interfering RNAs derived from dicer activity are typically about 21 to about 23 nucleotides in length and comprise about 19 base pair duplexes (Elbashir et al., 2001, Genes Dev., 15, 188). Dicer has also been implicated in the excision of 21-and 22-nucleotide small temporal RNAs (stRNAs) from precursor RNA of conserved structure that are implicated in translational control (Hutvagner et al., 2001, Science, 293, 834). The RNAi response also features an endonuclease complex, commonly referred to as an RNA-induced silencing complex (RISC), which mediates cleavage of single-stranded RNA having sequence complementary to the antisense strand of the siRNA duplex. Cleavage of the target RNA takes place in the middle of the region complementary to the antisense strand of the siRNA duplex. Cleavage of the target RNA takes place in the middle of the region complementary to the antisense strand of the siRNA duplex (Elbashir et al., 2001, Genes Dev., 15, 188).

RNAi has been studied in a variety of systems. Fire et al., 1998, Nature, 391, 806, were the first to observe RNAi in C. elegans. Wianny and Goetz, 1999, Nature Cell Biol., 2, 70, describe RNAi mediated by dsRNA in mouse embryos. Hammond et al., 2000, Nature, 404, 293, describe RNAi in Drosophila cells transfected with dsRNA. Elbashir et al., 2001, Nature, 411, 494, describe RNAi induced by introduction of duplexes of synthetic 21-nucleotide RNAs in cultured mammalian cells including human embryonic kidney and HeLa cells. Recent work in Drosophila embryonic lysates

5

10

15

20

25

(Elbashir et al., 2001, EMBO J., 20, 6877) has revealed certain requirements for siRNA length, structure, chemical composition, and sequence that are essential to mediate efficient RNAi activity. These studies have shown that 21-nucleotide siRNA duplexes are most active when containing 3'-terminal dinucleotide overhangs. Furthermore, complete substitution of one or both siRNA strands with 2'-deoxy (2'-H) or 2'-O-methyl nucleotides abolishes RNAi activity, whereas substitution of the 3'-terminal siRNA overhang nucleotides with 2'-deoxy nucleotides (2'-H) was shown to be tolerated. Single mismatch sequences in the center of the siRNA duplex were also shown to abolish RNAi activity. In addition, these studies also indicate that the position of the cleavage site in the target RNA is defined by the 5'-end of the siRNA guide sequence rather than the 3'-end of the guide sequence (Elbashir et al., 2001, EMBO J., 20, 6877). Other studies have indicated that a 5'-phosphate on the target-complementary strand of a siRNA duplex is required for siRNA activity and that ATP is utilized to maintain the 5'-phosphate moiety on the siRNA (Nykanen et al., 2001, Cell, 107, 309).

Studies have shown that replacing the 3'-terminal nucleotide overhanging segments -nucleotide duplex having two 3'-overhangs 21-mer siRNA deoxyribonucleotides does not have an adverse effect on RNAi activity. Replacing up to four nucleotides on each end of the siRNA with deoxyribonucleotides has been reported to be well tolerated, whereas complete substitution with deoxyribonucleotides results in no RNAi activity (Elbashir et al., 2001, EMBO J., 20, 6877). In addition, Elbashir et al., supra, also report that substitution of siRNA with 2'-O-methyl nucleotides completely abolishes RNAi activity. Li et al., International PCT Publication No. WO 00/44914, and Beach et al., International PCT Publication No. WO 01/68836 preliminarily suggest that siRNA may include modifications to either the phosphate-sugar backbone or the nucleoside to include at least one of a nitrogen or sulfur heteroatom, however, neither application postulates to what extent such modifications would be tolerated in siRNA molecules, nor provides any further guidance or examples of such modified siRNA. Kreutzer et al., Canadian Patent Application No. 2,359,180, also describe certain chemical modifications for use in dsRNA constructs in order to counteract activation of double-stranded RNA-dependent protein kinase PKR, specifically 2'-amino or 2'-Omethyl nucleotides, and nucleotides containing a 2'-O or 4'-C methylene bridge.

5

10

15

20

25

However, Kreutzer et al. similarly fails to provide examples or guidance as to what extent these modifications would be tolerated in siRNA molecules.

Parrish et al., 2000, Molecular Cell, 6, 1977-1087, tested certain chemical modifications targeting the unc-22 gene in C. elegans using long (>25 nt) siRNA transcripts. The authors describe the introduction of thiophosphate residues into these siRNA transcripts by incorporating thiophosphate nucleotide analogs with T7 and T3 RNA polymerase and observed that RNAs with two phosphorothicate modified bases also had substantial decreases in effectiveness as RNAi. Further, Parrish et al. reported that phosphorothicate modification of more than two residues greatly destabilized the RNAs in vitro such that interference activities could not be assayed. Id. at 1081. The authors also tested certain modifications at the 2'-position of the nucleotide sugar in the long siRNA transcripts and found that substituting deoxynucleotides for ribonucleotides produced a substantial decrease in interference activity, especially in the case of Uridine to Thymidine and/or Cytidine to deoxy-Cytidine substitutions. Id. In addition, the authors tested certain base modifications, including substituting, in sense and antisense strands of the siRNA, 4-thiouracil, 5-bromouracil, 5-iodouracil, and 3-(aminoallyl)uracil Whereas 4-thiouracil and 5-bromouracil for uracil, and inosine for guanosine. substitution appeared to be tolerated, Parrish reported that inosine produced a substantial decrease in interference activity when incorporated in either strand. Parrish also reported that incorporation of 5-iodouracil and 3-(aminoallyl)uracil in the antisense strand resulted in a substantial decrease in RNAi activity as well.

The use of longer dsRNA has been described. For example, Beach *et al.*, International PCT Publication No. WO 01/68836, describes specific methods for attenuating gene expression using endogenously-derived dsRNA. Tuschl *et al.*, International PCT Publication No. WO 01/75164, describe a *Drosophila in vitro* RNAi system and the use of specific siRNA molecules for certain functional genomic and certain therapeutic applications; although Tuschl, 2001, *Chem. Biochem.*, 2, 239-245, doubts that RNAi can be used to cure genetic diseases or viral infection due to the danger of activating interferon response. Li *et al.*, International PCT Publication No. WO 00/44914, describe the use of specific dsRNAs for attenuating the expression of certain target genes. Zernicka-Goetz *et al.*, International PCT Publication No. WO 01/36646, describe certain methods for inhibiting the expression of particular genes in mammalian

5

10

15

20

25

cells using certain dsRNA molecules. Fire et al., International PCT Publication No. WO 99/32619, describe particular methods for introducing certain dsRNA molecules into cells for use in inhibiting gene expression. Plaetinck et al., International PCT Publication No. WO 00/01846, describe certain methods for identifying specific genes responsible for conferring a particular phenotype in a cell using specific dsRNA molecules. Mello et al., International PCT Publication No. WO 01/29058, describe the identification of specific genes involved in dsRNA-mediated RNAi. Deschamps Depaillette et al., International PCT Publication No. WO 99/07409, describe specific compositions consisting of particular dsRNA molecules combined with certain anti-viral agents. Waterhouse et al., International PCT Publication No. 99/53050, describe certain methods for decreasing the phenotypic expression of a nucleic acid in plant cells using certain dsRNAs. Driscoll et al., International PCT Publication No. WO 01/49844, describe specific DNA constructs for use in facilitating gene silencing in targeted organisms.

Others have reported on various RNAi and gene-silencing systems. For example, Parrish et al., 2000, Molecular Cell, 6, 1977-1087, describe specific chemically-modified siRNA constructs targeting the unc-22 gene of C. elegans. Grossniklaus, International PCT Publication No. WO 01/38551, describes certain methods for regulating polycomb gene expression in plants using certain dsRNAs. Churikov et al., International PCT Publication No. WO 01/42443, describe certain methods for modifying genetic characteristics of an organism using certain dsRNAs. Cogoni et al., International PCT Publication No. WO 01/53475, describe certain methods for isolating a Neurospora silencing gene and uses thereof. Reed et al., International PCT Publication No. WO 01/68836, describe certain methods for gene silencing in plants. International PCT Publication No. WO 01/70944, describe certain methods of drug screening using transgenic nematodes as Parkinson's Disease models using certain dsRNAs. Deak et al., International PCT Publication No. WO 01/72774, describe certain Drosophila-derived gene products that may be related to RNAi in Drosophila. Arndt et al., International PCT Publication No. WO 01/92513 describe certain methods for mediating gene suppression by using factors that enhance RNAi. Tuschl et al., International PCT Publication No. WO 02/44321, describe certain synthetic siRNA Pachuk et al., International PCT Publication No. WO 00/63364, and Satishchandran et al., International PCT Publication No. WO 01/04313, describe certain

5

10

15

20

25

methods and compositions for inhibiting the function of certain polynucleotide sequences using certain dsRNAs. Echeverri *et al.*, International PCT Publication No. WO 02/38805, describe certain *C. elegans* genes identified via RNAi. Kreutzer *et al.*, International PCT Publications Nos. WO 02/055692, WO 02/055693, and EP 1144623 B1 describes certain methods for inhibiting gene expression using RNAi. Graham *et al.*, International PCT Publications Nos. WO 99/49029 and WO 01/70949, and AU 4037501 describe certain vector expressed siRNA molecules. Fire *et al.*, US 6,506,559, describe certain methods for inhibiting gene expression in vitro using certain long dsRNA (greater than 25 nucleotide) constructs that mediate RNAi.

## SUMMARY OF THE INVENTION

This invention relates to compounds, compositions, and methods useful for modulating RNA function and/or gene expression in a cell. Specifically, the instant invention features synthetic small nucleic acid molecules, such as short interfering nucleic acid (siNA), short interfering RNA (siRNA), double-stranded RNA (dsRNA), micro-RNA (miRNA), and short hairpin RNA (shRNA) molecules capable of modulating gene expression in cells by RNA inference (RNAi). The siRNA of the instant invention can be chemically synthesized, expressed from a vector or enzymatically synthesized. The use of chemically modified siNA can improve various properties of native siRNA molecules through increased resistance to nuclease degradation *in vivo* and/or improved cellular uptake. The chemically modified siNA molecules of the instant invention provide useful reagents and methods for a variety of therapeutic, diagnostic, agricultural, target validation, genomic discovery, genetic engineering and pharmacogenomic applications.

In a non-limiting example, the introduction of chemically modified nucleotides into nucleic acid molecules provides a powerful tool in overcoming potential limitations of *in vivo* stability and bioavailability inherent to native RNA molecules that are delivered exogenously. For example, the use of chemically modified nucleic acid molecules can enable a lower dose of a particular nucleic acid molecule for a given therapeutic effect since chemically modified nucleic acid molecules tend to have a longer half-life in serum. Furthermore, certain chemical modifications can improve the bioavailability of nucleic acid molecules by targeting particular cells or tissues and/or improving cellular uptake of the nucleic acid molecule. Therefore, even if the activity of a chemically modified

5

10

15

20

25

nucleic acid molecule is reduced as compared to a native nucleic acid molecule, for example when compared to an all RNA nucleic acid molecule, the overall activity of the modified nucleic acid molecule can be greater than the native molecule due to improved stability and/or delivery of the molecule. Unlike native unmodified siRNA, chemically modified siNA can also minimize the possibility of activating interferon activity in humans.

The siRNA molecules of the invention can be designed to inhibit gene expression through RNAi targeting of a variety of RNA molecules. In one embodiment, the siRNA molecules of the invention are used to target various RNAs corresponding to a target gene. Non-limiting examples of such RNAs include messenger RNA (mRNA), alternate RNA splice variants of target gene(s), post-transcriptionally modified RNA of target gene(s), pre-mRNA of target gene(s). If alternate splicing produces a family of transcipts that are distinguished by usage of appropriate exons, the instant invention can be used to inhibit gene expression through the appropriate exons to specifically inhibit or to distinguish among the functions of gene family members. For example, a protein that contains an alternatively spliced transmembrane domain can be expressed in both membrane bound and secreted forms. Use of the invention to target the exon containing the transmembrane domain can be used to determine the functional consequences of pharmaceutical targeting of membrane bound as opposed to the secreted form of the protein. Non-limiting examples of applications of the invention relating to targeting these RNA molecules include therapeutic pharmaceutical applications, pharmaceutical discovery applications, molecular diagnostic and gene function applications, and gene mapping, for example using single nucleotide polymorphism mapping with siRNA molecules of the invention. Such applications can be implemented using known gene sequences or from partial sequences available from an expressed sequence tag (EST).

In another embodiment, the siRNA molecules of the invention are used to target conserved sequences corresponding to a gene family or gene families. As such, siRNA can be used to characterize pathways of gene function in a variety of applications. For example, the present invention can be used to inhibit the activity of target gene(s) in a pathway to determine the function of uncharacterized gene(s) in gene function analysis, mRNA function analysis, or translational analysis. The invention can be used to determine potential target gene pathways involved in various diseases and conditions

5

10

15

20

25

toward pharmaceutical development. The invention can be used to understand pathways of gene expression involved in development, such as prenatal development, postnatal development and/or aging.

In one embodiment, the invention features a short interfering nucleic acid (siNA) molecule that down-regulates expression of a gene family by RNA interference. The gene family can comprise more than one splice variant of a target gene, more than one post-transcriptionally modified RNA of a target gene, or more than one RNA trascript having shared homology. In one embodiment, the gene family comprises epidermal growth factor (e.g., EGFR, such as HER1, HER2, HER3, and/or HER4) genes, vascular endothelial growth factor and vascular endothelial growth factor receptor (e.g., VEGF, VEGFR1, VEGFR2, or VEGFR3) genes, or viral genes corresponding to different viral strains (e.g., HIV-1 and HIV-2). Such gene families can be established by analysing nucleic acid sequences (e.g., sequences shown by Genbank Accession Nos. in Table V) for homology.

In one embodiment, the invention features a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene (e.g., a human gene), wherein the siNA molecule comprises one or more chemical modifications and each strand of the double-stranded siNA is about 21 nucleotides long.

In one embodiment, a siNA molecule of the invention comprises no ribonucleotides. In another embodiment, a siNA molecule of the invention comprises ribonucleotides.

In one embodiment, the invention features a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene (e.g., a human gene), wherein one of the strands of the double-stranded siNA molecule comprises a nucleotide sequence that is complementary to a nucleotide sequence of the endogenous mammalian target gene or a portion thereof, and wherein the second strand of the double-stranded siNA molecule comprises a nucleotide sequence substantially similar to the nucleotide sequence of the endogenous mammalian target gene or a portion thereof.

5

10

15

20

25

In one embodiment, the invention features a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene (e.g., a human gene), wherein each strand of the siNA molecule comprises about 19 to about 23 nucleotides, and wherein each strand comprises about 19 nucleotides that are complementary to the nucleotides of the other strand.

In one embodiment, the invention features a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene (e.g., a human gene), wherein the siNA molecule comprises an antisense region comprising a nucleotide sequence that is complementary to a nucleotide sequence of the endogenous mammalian target gene or a portion thereof, and wherein the siNA further comprises a sense region, wherein the sense region comprises a nucleotide sequence substantially similar to the nucleotide sequence of the endogenous mammalian target gene or a portion thereof.

In one embodiment, the invention features a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene (e.g., a human gene), wherein the antisense region and the sense region each comprise about 19 to about 23 nucleotides, and wherein the antisense region comprises about 19 nucleotides that are complementary to nucleotides of the sense region.

In one embodiment, the invention features a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene (e.g., a human gene), wherein the siNA molecule comprises a sense region and an antisense region and wherein the antisense region comprises a nucleotide sequence that is complementary to a nucleotide sequence of RNA encoded by the endogenous mammalian target gene or a portion thereof and the sense region comprises a nucleotide sequence that is complementary to the antisense region.

In one embodiment, the invention features a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene (e.g., a human gene), wherein the siNA molecule is assembled from two separate oligonucleotide fragments wherein one fragment comprises the sense region and the second fragment comprises the antisense region of the siNA molecule.

5

10

15

20

25

The sense region can be connected to the antisense region via a linker molecule, such as a polynucleotide linker or a non-nucleotide linker.

In one embodiment, the invention features a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene (e.g., a human gene), wherein the siNA molecule comprises a sense region and an antisense region and wherein the antisense region comprises a nucleotide sequence that is complementary to a nucleotide sequence of RNA encoded by the endogenous mammalian target gene or a portion thereof and the sense region comprises a nucleotide sequence that is complementary to the antisense region, and wherein pyrimidine nucleotides in the sense region are 2'-O-methyl pyrimidine nucleotides, 2'-deoxy nucleotides, and/or 2'-deoxy-2'-fluoro pyrimidine nucleotides.

5

10

15

20

25

30

BNSDOCID: <WO\_

03074654A2\_I\_>

In one embodiment, the invention features a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene (e.g., a human gene), wherein the siNA molecule is assembled from two separate oligonucleotide fragments wherein one fragment comprises the sense region and the second fragment comprises the antisense region of the siNA molecule, and wherein the fragment comprising the sense region includes a terminal cap moiety at the 5'-end, the 3'-end, or both of the 5' and 3' ends of the fragment comprising the sense region. In another embodiment, the terminal cap moiety is an inverted deoxy abasic moiety or glyceryl moiety. In another embodiment, each of the two fragments of the siNA molecule comprise 21 nucleotides.

In one embodiment, the invention features a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene (e.g., a human gene), wherein the siNA molecule comprises a sense region and an antisense region and wherein the antisense region comprises a nucleotide sequence that is complementary to a nucleotide sequence of RNA encoded by the endogenous mammalian target gene or a portion thereof and the sense region comprises a nucleotide sequence that is complementary to the antisense region, and wherein the purine nucleotides present in the antisense region comprise 2'-deoxy- purine nucleotides. In another embodiment, the antisense region comprises a phosphorothioate

internucleotide linkage at the 3' end of the antisense region. In another embodiment, the antisense region comprises a glyceryl modification at the 3' end of the antisense region.

In one embodiment, the invention features a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene (e.g., a human gene), wherein the siNA molecule is assembled from two separate oligonucleotide fragments wherein one fragment comprises the sense region and the second fragment comprises the antisense region of the siNA molecule, and wherein about 19 nucleotides of each fragment of the siNA molecule are base-paired to the complementary nucleotides of the other fragment of the siNA molecule and wherein at least two 3' terminal nucleotides of each fragment of the siNA molecule are not basepaired to the nucleotides of the other fragment of the siNA molecule. In another embodiment, each of the two 3' terminal nucleotides of each fragment of the siNA molecule are 2'-deoxy-pyrimidines, such as 2'-deoxy-thymidine. In another embodiment, all 21 nucleotides of each fragment of the siNA molecule are base-paired to the complementary nucleotides of the other fragment of the siNA molecule. In another embodiment, about 19 nucleotides of the antisense region are base-paired to the nucleotide sequence or a portion thereof of the RNA encoded by the endogenous mammalian target gene. In another embodiment, 21 nucleotides of the antisense region are base-paired to the nucleotide sequence or a portion thereof of the RNA encoded by the endogenous mammalian target gene. In another embodiment, the 5'-end of the fragment comprising said antisense region optionally includes a phosphate group.

In one embodiment, the invention features a double-stranded short interfering nucleic acid (siNA) molecule that inhibits the expression of an endogenous mammalian target RNA sequence (e.g., wherein said target RNA sequence is encoded by a human gene), wherein the siNA molecule comprises no ribonucleotides and wherein each strand of the double-stranded siNA molecule comprises about 21 nucleotides.

In one embodiment, the invention features a double-stranded short interfering nucleic acid (siNA) molecule that inhibits the expression of an endogenous mammalian target gene (e.g., a human gene such as vascular endothelial growth factor, vascular endothelial growth factor receptor (such as VEGFR1, VEGFR2, or VEGFR3), BCL2, HER2/neu, c-Myc, PCNA, REL-A, PTP1B, BACE, CHK1, PKC-alpha, or EGFR).

5

10

15

20

25

wherein the siNA molecule does not require the presence of a ribonucleotide within the siNA molecule for said inhibition of expression of an endogenous mammalian target gene and wherein each strand of the double-stranded siNA molecule is about 21 nucleotides long.

In one embodiment, the invention features a medicament comprising a siNA molecule of the invention.

In one embodiment, the invention features an active ingredient comprising a siNA molecule of the invention.

In one embodiment, the invention features the use of a double-stranded short interfering nucleic acid (siNA) molecule to down-regulate expression of an endogenous mammalian target gene, wherein the siNA molecule comprises one or more chemical modifications and each strand of the double-stranded siNA is about 21 nucleotides long.

In one embodiment, siRNA molecule(s) and/or methods of the invention are used to inhibit the expression of gene(s) that encode RNA referred to by Genbank Accession number in **Table V**. In another embodiment, siRNA molecule(s) and/or methods of the invention are used to target RNA sequence(s) referred to by Genbank Accession number in **Table V**, or nucleic acid sequences encoding such sequences referred to by Genbank Accession number in **Table V**. Such sequences are readily obtained using the Genbank Accession numbers in **Table V**.

In one embodiment, the invention features a siNA molecule having RNAi activity against an RNA encoding a protein, wherein the siNA molecule comprises a sequence complementary to RNA having protein encoding sequence, such as those sequences having GenBank Accession Nos. shown in **Table V**.

In another embodiment, the invention features a siNA molecule having RNAi activity against a gene, wherein the siNA molecule comprises nucleotide sequence complementary to a nucleotide sequence of the gene, such as genes encoding sequences having GenBank Accession Nos. shown in Table V. In another embodiment, a siNA molecule of the invention includes nucleotide sequence that can interact with nucleotide sequence of a gene and thereby mediate silencing of gene expression, for example,

5

10

15

wherein the siNA mediates regulation of gene expression by cellular processes that modulate the chromatin structure of the gene and prevent transcription of the gene.

In yet another embodiment, the invention features a siNA molecule comprising a sequence, for example, the antisense sequence of the siNA construct, complementary to a sequence represented by GenBank Accession Nos. shown in **Table V** or a portion of said sequence.

In one embodiment, the nucleic acid molecules of the invention that act as mediators of the RNA interference gene silencing response are chemically modified double stranded nucleic acid molecules. As in their native double stranded RNA counterparts, these siNA molecules typically consist of duplexes containing about 19 base pairs between oligonucleotides comprising about 19 to about 25 nucleotides. The most active siRNA molecules are thought to have such duplexes with overhanging ends of 1-3 nucleotides, for example 21 nucleotide duplexes with 19 base pairs and 2 nucleotide 3'overhangs. These overhanging segments are readily hydrolyzed by endonucleases in vivo. Studies have shown that replacing the 3'-overhanging segments of a 21-mer siRNA duplex having 2 nucleotide 3' overhangs with deoxyribonucleotides does not have an adverse effect on RNAi activity. Replacing up to 4 nucleotides on each end of the siRNA with deoxyribonucleotides has been reported to be well tolerated whereas complete substitution with deoxyribonucleotides results in no RNAi activity (Elbashir et al., 2001, EMBO J., 20, 6877). In addition, Elbashir et al, supra, also report that substitution of siRNA with 2'-O-methyl nucleotides completely abolishes RNAi activity. Li et al., International PCT Publication No. WO 00/44914, and Beach et al., International PCT Publication No. WO 01/68836 both suggest that siRNA may include modifications to either the phosphate-sugar back bone or the nucleoside to include at least one of a nitrogen or sulfur heteroatom, however neither application teaches to what extent these modifications are tolerated in siRNA molecules nor provide any examples of such modified siRNA. Kreutzer and Limmer, Canadian Patent Application No. 2,359,180, also describe certain chemical modifications for use in dsRNA constructs in order to counteract activation of double stranded-RNA-dependent protein kinase PKR, specifically 2'-amino or 2'-O-methyl nucleotides, and nucleotides containing a 2'-O or 4'-C methylene bridge. However, Kreutzer and Limmer similarly fail to show to what

5

10

15

20

25

extent these modifications are tolerated in siRNA molecules nor provide any examples of such modified siRNA.

In one embodiment, the invention features chemically modified siNA constructs having specificity for target nucleic acid molecules in a cell (i.e. target nucleic acid molecules comprising or encoded by sequences referred to herein by Genbank Accession numbers in Table V). Non-limiting examples of such chemical modifications include linkages, 2'-O-methyl internucleotide phosphorothioate without limitation ribonucleotides, 2'-deoxy-2'-fluoro ribonucleotides, 2'-deoxy ribonucleotides, "universal base" nucleotides, 5-C-methyl nucleotides, and inverted deoxyabasic residue incorporation. These chemical modifications, when used in various siNA constructs, are shown to preserve RNAi activity in cells while at the same time, dramatically increasing the serum stability of these compounds. Furthermore, contrary to the data published by Parrish et al., supra, applicant demonstrates that multiple (greater than one) phosphorothioate substitutions are well-tolerated and confer substantial increases in serum stability for modified siNA constructs.

In one embodiment, a siNA molecule of the invention comprises modified nucleotides while maintaining the ability to mediate RNAi. The modified nucleotides can be used to improve in vitro or in vivo characteristics such as stability, activity, and/or bioavailability. For example, a siNA molecule of the invention can comprise modified nucleotides as a percentage of the total number of nucleotides present in the siNA molecule. As such, a siNA molecule of the invention can generally comprise modified nucleotides of about 5 to about 100% of the nucleotide positions (e.g., 5%, 10%, 15%, 20%, 25%, 30%, 35%, 40%, 45%, 50%, 55%, 60%, 65%, 70%, 75%, 80%, 85%, 90%, 95% or 100% of the nucleotide positions). The actual percentage of modified nucleotides present in a given siNA molecule depends on the total number of nucleotides present in the siNA. If the siNA molecule is single stranded, the percent modification can be based upon the total number of nucleotides present in the single stranded siNA molecules. Likewise, if the siNA molecule is double stranded, the percent modification can be based upon the total number of nucleotides present in the sense strand, antisense strand, or both In addition, the actual percentage of modified the sense and antisense strands. nucleotides present in a given siNA molecule can also depend on the total number of purine and pyrimidine nucleotides present in the siNA, for example wherein all

5

10

15

20

25

pyrimidine nucleotides and/or all purine nucleotides present in the siNA molecule are modified.

The antisense region of a siNA molecule of the invention can comprise a phosphorothioate internucleotide linkage at the 3'-end of said antisense region. The antisense region can comprise about one to about five phosphorothioate internucleotide linkages at the 5'-end of said antisense region. The 3'-terminal nucleotide overhangs of a siNA molecule of the invention can comprise ribonucleotides or deoxyribonucleotides that are chemically-modified at a nucleic acid sugar, base, or backbone. The 3'-terminal nucleotide overhangs can comprise one or more universal base ribonucleotides. The 3'-terminal nucleotide overhangs can comprise one or more acyclic nucleotides.

5

10

15

20

25

BNSDOCID: <WO\_

03074654A2 I >

One embodiment of the invention provides an expression vector comprising a nucleic acid sequence encoding at least one siNA molecule of the invention in a manner that allows expression of the nucleic acid molecule. Another embodiment of the invention provides a mammalian cell comprising such an expression vector. The mammalian cell can be a human cell. The siNA molecule of the expression vector can comprise a sense region and an antisense region. The antisense region can comprise sequence complementary to a RNA or DNA sequence encoding a protein and the sense region can comprise sequence complementary to the antisense region. The siNA molecule can comprise two distinct strands having complementary sense and antisense regions. The siNA molecule can comprise a single strand having complementary sense and antisense regions.

In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule capable of mediating RNA interference (RNAi) inside a cell or reconstituted *in vitro* system, wherein the chemical modification comprises one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) nucleotides comprising a backbone modified internucleotide linkage having Formula I:

$$R_1$$
— $X$ — $P$ — $Y$ — $R_2$ 

wherein each R1 and R2 is independently any nucleotide, non-nucleotide, or polynucleotide which can be naturally-occurring or chemically-modified, each X and Y is independently O, S, N, alkyl, or substituted alkyl, each Z and W is independently O, S, N, alkyl, substituted alkyl, O-alkyl, S-alkyl, alkaryl, or aralkyl, and wherein W, X, Y, and Z are optionally not all O.

5

10

15

20

25

30

BNSDOCID: <WO\_\_\_\_\_

\_03074654A2\_l\_>

The chemically-modified internucleotide linkages having Formula I, for example, wherein any Z, W, X, and/or Y independently comprises a sulphur atom, can be present in one or both oligonucleotide strands of the siNA duplex, for example, in the sense strand, the antisense strand, or both strands. The siNA molecules of the invention can comprise one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) chemicallymodified internucleotide linkages having Formula I at the 3'-end, the 5'-end, or both of the 3' and 5'-ends of the sense strand, the antisense strand, or both strands. For example, an exemplary siNA molecule of the invention can comprise about 1 to about 5 or more (e.g., about 1, 2, 3, 4, 5, or more) chemically-modified internucleotide linkages having Formula I at the 5'-end of the sense strand, the antisense strand, or both strands. In another non-limiting example, an exemplary siNA molecule of the invention can comprise one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) pyrimidine nucleotides with chemically-modified internucleotide linkages having Formula I in the sense strand, the antisense strand, or both strands. In yet another non-limiting example, an exemplary siNA molecule of the invention can comprise one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) purine nucleotides with chemically-modified internucleotide linkages having Formula I in the sense strand, the antisense strand, or both strands. In another embodiment, a siNA molecule of the invention having internucleotide linkage(s) of Formula I also comprises a chemically-modified nucleotide or nonnucleotide having any of Formulae I-VII.

In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule capable of mediating RNA interference (RNAi) inside a cell or reconstituted *in vitro* system, wherein the chemical modification comprises one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) nucleotides or non-nucleotides having Formula II:

wherein each R3, R4, R5, R6, R7, R8, R10, R11 and R12 is independently H, OH, alkyl, substituted alkyl, alkaryl or aralkyl, F, Cl, Br, CN, CF3, OCF3, OCN, O-alkyl, S-alkyl, N-alkyl, O-alkyl, S-alkyl, N-alkyl, O-alkyl-OH, O-alkyl-OH, O-alkyl-OH, S-alkyl-SH, alkyl-S-alkyl, alkyl-O-alkyl, ONO2, NO2, N3, NH2, aminoalkyl, aminoacid, aminoacyl, ONH2, O-aminoalkyl, O-aminoacid, O-aminoacyl, heterocycloalkyl, heterocycloalkaryl, aminoalkylamino, polyalklylamino, substituted silyl, or group having Formula I; R9 is O, S, CH2, S=O, CHF, or CF2, and B is a nucleosidic base such as adenine, guanine, uracil, cytosine, thymine, 2-aminoadenosine, 5-methylcytosine, 2,6-diaminopurine, or any other non-naturally occurring base that can be complementary or non-complementary to target RNA or a non-nucleosidic base such as phenyl, naphthyl, 3-nitropyrrole, 5-nitroindole, nebularine, pyridone, pyridinone, or any other non-naturally occurring universal base that can be complementary or non-complementary to target RNA.

5

10

15

20

25

BNSDOCID: <WO\_\_\_

03074654A2\_l\_>

The chemically-modified nucleotide or non-nucleotide of Formula II can be present in one or both oligonucleotide strands of the siNA duplex, for example in the sense strand, the antisense strand, or both strands. The siNA molecules of the invention can comprise one or more chemically-modified nucleotide or non-nucleotide of Formula II at the 3'-end, the 5'-end, or both of the 3' and 5'-ends of the sense strand, the antisense strand, or both strands. For example, an exemplary siNA molecule of the invention can comprise about 1 to about 5 or more (e.g., about 1, 2, 3, 4, 5, or more) chemically-modified nucleotides or non-nucleotides of Formula II at the 5'-end of the sense strand, the antisense strand, or both strands. In anther non-limiting example, an exemplary siNA molecule of the invention can comprise about 1 to about 5 or more (e.g., about 1, 2, 3, 4, 5, or more) chemically-modified nucleotides or non-nucleotides of Formula II at the 3'-end of the sense strand, the antisense strand, or both strands.

In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule capable of mediating RNA interference (RNAi) inside a cell or reconstituted *in vitro* system, wherein the chemical modification comprises one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) nucleotides or non-nucleotides having Formula III:

5

10

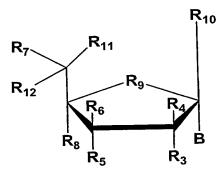
15

20

25

BNSDOCID: <WO

03074654A2 1\_>



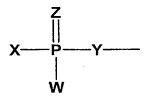
wherein each R3, R4, R5, R6, R7, R8, R10, R11 and R12 is independently H, OH, alkyl, substituted alkyl, alkaryl or aralkyl, F, Cl, Br, CN, CF3, OCF3, OCN, O-alkyl, S-alkyl, N-alkyl, O-alkyl, S-alkyl, N-alkyl, O-alkyl, S-alkyl, S-alkyl, S-alkyl-OH, S-alkyl-SH, S-alkyl-SH, alkyl-O-alkyl, ONO2, NO2, N3, NH2, aminoalkyl, aminoacid, aminoacyl, ONH2, O-aminoalkyl, O-aminoacid, O-aminoacyl, heterocycloalkyl, heterocycloalkaryl, aminoalkylamino, polyalklylamino, substituted silyl, or group having Formula I; R9 is O, S, CH2, S=O, CHF, or CF2, and B is a nucleosidic base such as adenine, guanine, uracil, cytosine, thymine, 2-aminoadenosine, 5-methylcytosine, 2,6-diaminopurine, or any other non-naturally occurring base that can be employed to be complementary or non-complementary to target RNA or a non-nucleosidic base such as phenyl, naphthyl, 3-nitropyrrole, 5-nitroindole, nebularine, pyridone, pyridinone, or any other non-naturally occurring universal base that can be complementary or non-complementary to target RNA.

The chemically-modified nucleotide or non-nucleotide of Formula III can be present in one or both oligonucleotide strands of the siNA duplex, for example, in the sense strand, the antisense strand, or both strands. The siNA molecules of the invention can comprise one or more chemically-modified nucleotide or non-nucleotide of Formula III at the 3'-end, the 5'-end, or both of the 3' and 5'-ends of the sense strand, the antisense strand, or both strands. For example, an exemplary siNA molecule of the invention can comprise about 1 to about 5 or more (e.g., about 1, 2, 3, 4, 5, or more) chemically-

modified nucleotide(s) or non-nucleotide(s) of Formula III at the 5'-end of the sense strand, the antisense strand, or both strands. In anther non-limiting example, an exemplary siNA molecule of the invention can comprise about 1 to about 5 or more (e.g., about 1, 2, 3, 4, 5, or more) chemically-modified nucleotide or non-nucleotide of Formula III at the 3'-end of the sense strand, the antisense strand, or both strands.

In another embodiment, a siNA molecule of the invention comprises a nucleotide having Formula II or III, wherein the nucleotide having Formula II or III is in an inverted configuration. For example, the nucleotide having Formula II or III is connected to the siNA construct in a 3'-3', 3'-2', 2'-3', or 5'-5' configuration, such as at the 3'-end, the 5'-end, or both of the 3' and 5'-ends of one or both siNA strands.

In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule capable of mediating RNA interference (RNAi) inside a cell or reconstituted *in vitro* system, wherein the chemical modification comprises a 5'-terminal phosphate group having Formula IV:



15

20

25

5

10

wherein each X and Y is independently O, S, N, alkyl, substituted alkyl, or alkylhalo; wherein each Z and W is independently O, S, N, alkyl, substituted alkyl, O-alkyl, S-alkyl, alkaryl, aralkyl, or alkylhalo; and wherein W, X, Y and Z are not all O.

In one embodiment, the invention features a siNA molecule having a 5'-terminal phosphate group having Formula IV on the target-complementary strand, for example, a strand complementary to a target RNA, wherein the siNA molecule comprises an all RNA siNA molecule. In another embodiment, the invention features a siNA molecule having a 5'-terminal phosphate group having Formula IV on the target-complementary strand wherein the siNA molecule also comprises about 1 to about 3 (e.g., about 1, 2, or 3) nucleotide 3'-terminal nucleotide overhangs having about 1 to about 4 (e.g., about 1, 2, 3, or 4) deoxyribonucleotides on the 3'-end of one or both strands. In another embodiment, a 5'-terminal phosphate group having Formula IV is present on the target-complementary

strand of a siNA molecule of the invention, for example a siNA molecule having chemical modifications having any of Formulae I-VII.

In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule capable of mediating RNA interference (RNAi) inside a cell or reconstituted in vitro system, wherein the chemical modification comprises one or more phosphorothioate internucleotide linkages. For example, in a non-limiting example, the invention features a chemically-modified short interfering nucleic acid (siNA) having about 1, 2, 3, 4, 5, 6, 7, 8 or more phosphorothioate internucleotide linkages in one siNA strand. In yet another embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) individually having about 1, 2, 3, 4, 5, 6, 7, 8 or more phosphorothioate internucleotide linkages in both siNA strands. The phosphorothioate internucleotide linkages can be present in one or both oligonucleotide strands of the siNA duplex, for example in the sense strand, the antisense strand, or both strands. The siNA molecules of the invention can comprise one or more phosphorothioate internucleotide linkages at the 3'-end, the 5'-end, or both of the 3'- and 5'-ends of the sense strand, the antisense strand, or both strands. For example, an exemplary siNA molecule of the invention can comprise about 1 to about 5 or more (e.g., about 1, 2, 3, 4, 5, or more) consecutive phosphorothioate internucleotide linkages at the 5'-end of the sense strand, the antisense strand, or both strands. In another non-limiting example, an exemplary siNA molecule of the invention can comprise one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) pyrimidine phosphorothioate internucleotide linkages in the sense strand, the antisense strand, or both strands. In yet another non-limiting example, an exemplary siNA molecule of the invention can comprise one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) purine phosphorothioate internucleotide linkages in the sense strand, the antisense strand, or both strands.

In one embodiment, the invention features a siNA molecule, wherein the sense strand comprises one or more, for example, about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more phosphorothicate internucleotide linkages, and/or one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more) 2'-deoxy, 2'-O-methyl, 2'-deoxy-2'-fluoro, and/or about one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more) universal base modified nucleotides, and optionally a terminal cap molecule at the 3'-end, the 5'-end, or both of the 3'- and 5'-ends of the sense strand; and wherein the antisense strand comprises about 1 to about 10 or

5

10

15

20

25

more, specifically about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more phosphorothioate internucleotide linkages, and/or one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more) 2'-deoxy, 2'-O-methyl, 2'-deoxy-2'-fluoro, and/or one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more) universal base modified nucleotides, and optionally a terminal cap molecule at the 3'-end, the 5'-end, or both of the 3'- and 5'-ends of the antisense strand. In another embodiment, one or more, for example about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more, pyrimidine nucleotides of the sense and/or antisense siNA strand are chemically-modified with 2'-deoxy, 2'-O-methyl and/or 2'-deoxy-2'-fluoro nucleotides, with or without one or more, for example about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more, phosphorothioate internucleotide linkages and/or a terminal cap molecule at the 3'-end, the 5'-end, or both of the 3'- and 5'-ends, being present in the same or different strand.

In another embodiment, the invention features a siNA molecule, wherein the sense strand comprises about 1 to about 5, specifically about 1, 2, 3, 4, or 5 phosphorothioate internucleotide linkages, and/or one or more (e.g., about 1, 2, 3, 4, 5, or more) 2'-deoxy, 2'-O-methyl, 2'-deoxy-2'-fluoro, and/or one or more (e.g., about 1, 2, 3, 4, 5, or more) universal base modified nucleotides, and optionally a terminal cap molecule at the 3-end, the 5'-end, or both of the 3'- and 5'-ends of the sense strand; and wherein the antisense strand comprises about 1 to about 5 or more, specifically about 1, 2, 3, 4, 5, or more phosphorothioate internucleotide linkages, and/or one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more) 2'-deoxy, 2'-O-methyl, 2'-deoxy-2'-fluoro, and/or one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more) universal base modified nucleotides, and optionally a terminal cap molecule at the 3'-end, the 5'-end, or both of the 3'- and 5'-ends of the antisense strand. In another embodiment, one or more, for example about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more, pyrimidine nucleotides of the sense and/or antisense siNA strand are chemically-modified with 2'-deoxy, 2'-O-methyl and/or 2'-deoxy-2'-fluoro nucleotides, with or without about 1 to about 5 or more, for example about 1, 2, 3, 4, 5, or more phosphorothioate internucleotide linkages and/or a terminal cap molecule at the 3'end, the 5'-end, or both of the 3'- and 5'-ends, being present in the same or different strand.

In one embodiment, the invention features a siNA molecule, wherein the antisense strand comprises one or more, for example, about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more phosphorothioate internucleotide linkages, and/or about one or more (e.g., about 1, 2, 3,

5

10

15

20

25

4, 5, 6, 7, 8, 9, 10 or more) 2'-deoxy, 2'-O-methyl, 2'-deoxy-2'-fluoro, and/or one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more) universal base modified nucleotides, and optionally a terminal cap molecule at the 3'-end, the 5'-end, or both of the 3'- and 5'-ends of the sense strand; and wherein the antisense strand comprises about 1 to about 10 or more, specifically about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more phosphorothioate internucleotide linkages, and/or one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more) 2'-deoxy, 2'-O-methyl, 2'-deoxy-2'-fluoro, and/or one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more) universal base modified nucleotides, and optionally a terminal cap molecule at the 3'-end, the 5'-end, or both of the 3'- and 5'-ends of the antisense strand. In another embodiment, one or more, for example about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more pyrimidine nucleotides of the sense and/or antisense siNA strand are chemically-modified with 2'-deoxy, 2'-O-methyl and/or 2'-deoxy-2'-fluoro nucleotides, with or without one or more, for example, about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more phosphorothioate internucleotide linkages and/or a terminal cap molecule at the 3'-end, the 5'-end, or both of the 3' and 5'-ends, being present in the same or different strand.

5

10

15

20

25

30

BNSDOCID: <WO\_

03074654A2\_1\_>

In another embodiment, the invention features a siNA molecule, wherein the antisense strand comprises about 1 to about 5 or more, specifically about 1, 2, 3, 4, 5 or more phosphorothioate internucleotide linkages, and/or one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more) 2'-deoxy, 2'-O-methyl, 2'-deoxy-2'-fluoro, and/or one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more) universal base modified nucleotides, and optionally a terminal cap molecule at the 3'-end, the 5'-end, or both of the 3'- and 5'-ends of the sense strand; and wherein the antisense strand comprises about 1 to about 5 or more, specifically about 1, 2, 3, 4, 5 or more phosphorothioate internucleotide linkages, and/or one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more) 2'-deoxy, 2'-O-methyl, 2'-deoxy-2'-fluoro, and/or one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more) universal base modified nucleotides, and optionally a terminal cap molecule at the 3'-end, the 5'-end, or both of the 3'- and 5'-ends of the antisense strand. In another embodiment, one or more, for example about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more pyrimidine nucleotides of the sense and/or antisense siNA strand are chemically-modified with 2'-deoxy, 2'-Omethyl and/or 2'-deoxy-2'-fluoro nucleotides, with or without about 1 to about 5, for example about 1, 2, 3, 4, 5 or more phosphorothioate internucleotide linkages and/or a

terminal cap molecule at the 3'-end, the 5'-end, or both of the 3'- and 5'-ends, being present in the same or different strand.

In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule having about 1 to about 5, specifically about 1, 2, 3, 4, 5 or more phosphorothioate internucleotide linkages in each strand of the siNA molecule.

In another embodiment, the invention features a siNA molecule comprising 2'-5' internucleotide linkages. The 2'-5' internucleotide linkage(s) can be at the 3'-end, the 5'-end, or both of the 3'- and 5'-ends of one or both siNA sequence strands. In addition, the 2'-5' internucleotide linkage(s) can be present at various other positions within one or both siNA sequence strands, for example, about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more including every internucleotide linkage of a pyrimidine nucleotide in one or both strands of the siNA molecule can comprise a 2'-5' internucleotide linkage, or about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more including every internucleotide linkage of a purine nucleotide in one or both strands of the siNA molecule can comprise a 2'-5' internucleotide linkage.

In another embodiment, a chemically-modified siNA molecule of the invention comprises a duplex having two strands, one or both of which can be chemically-modified, wherein each strand is about 18 to about 27 (e.g., about 18, 19, 20, 21, 22, 23, 24, 25, 26, or 27) nucleotides in length, wherein the duplex has about 18 to about 23 (e.g., about 18, 19, 20, 21, 22, or 23) base pairs, and wherein the chemical modification comprises a structure having any of Formulae I-VII. For example, an exemplary chemically-modified siNA molecule of the invention comprises a duplex having two strands, one or both of which can be chemically-modified with a chemical modification having any of Formulae I-VII or any combination thereof, wherein each strand consists of about 21 nucleotides, each having a 2-nucleotide 3'-terminal nucleotide overhang, and wherein the duplex has about 19 base pairs. In another embodiment, a siNA molecule of the invention comprises a single stranded hairpin structure, wherein the siNA is about 36 to about 70 (e.g., about 36, 40, 45, 50, 55, 60, 65, or 70) nucleotides in length having about 18 to about 23 (e.g., about 18, 19, 20, 21, 22, or 23) base pairs, and wherein the siNA can include a chemical modification comprising a structure having any of Formulae I-VII or any combination thereof. For example, an exemplary chemically-modified siNA molecule of the invention comprises a linear oligonucleotide having about 42 to about 50 (e.g., about 42, 43, 44, 45,

5

10

15

20

25

46, 47, 48, 49, or 50) nucleotides that is chemically-modified with a chemical modification having any of Formulae I-VII or any combination thereof, wherein the linear oligonucleotide forms a hairpin structure having about 19 base pairs and a 2-nucleotide 3'-terminal nucleotide overhang. In another embodiment, a linear hairpin siNA molecule of the invention contains a stem loop motif, wherein the loop portion of the siNA molecule is biodegradable. For example, a linear hairpin siNA molecule of the invention is designed such that degradation of the loop portion of the siNA molecule *in vivo* can generate a double-stranded siNA molecule with 3'-terminal overhangs, such as 3'-terminal nucleotide overhangs comprising about 2 nucleotides.

In another embodiment, a siNA molecule of the invention comprises a circular nucleic acid molecule, wherein the siNA is about 38 to about 70 (e.g., about 38, 40, 45, 50, 55, 60, 65, or 70) nucleotides in length having about 18 to about 23 (e.g., about 18, 19, 20, 21, 22, or 23) base pairs, and wherein the siNA can include a chemical modification, which comprises a structure having any of Formulae I-VII or any combination thereof. For example, an exemplary chemically-modified siNA molecule of the invention comprises a circular oligonucleotide having about 42 to about 50 (e.g., about 42, 43, 44, 45, 46, 47, 48, 49, or 50) nucleotides that is chemically-modified with a chemical modification having any of Formulae I-VII or any combination thereof, wherein the circular oligonucleotide forms a dumbbell shaped structure having about 19 base pairs and 2 loops.

In another embodiment, a circular siNA molecule of the invention contains two loop motifs, wherein one or both loop portions of the siNA molecule is biodegradable. For example, a circular siNA molecule of the invention is designed such that degradation of the loop portions of the siNA molecule *in vivo* can generate a double-stranded siNA molecule with 3'-terminal overhangs, such as 3'-terminal nucleotide overhangs comprising about 2 nucleotides.

In one embodiment, a siNA molecule of the invention comprises at least one (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) abasic moiety, for example a compound having Formula V:

5

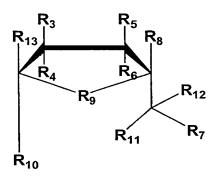
10

15

20

wherein each R3, R4, R5, R6, R7, R8, R10, R11, R12, and R13 is independently H, OH, alkyl, substituted alkyl, alkaryl or aralkyl, F, Cl, Br, CN, CF3, OCF3, OCN, O-alkyl, S-alkyl, N-alkyl, O-alkyl, S-alkyl, N-alkyl, S-alkyl, N-alkyl-OSH, alkyl-OSH, alkyl-OH, O-alkyl-OH, O-alkyl-SH, S-alkyl-OH, S-alkyl-SH, alkyl-S-alkyl, alkyl-O-alkyl, ONO2, NO2, N3, NH2, aminoalkyl, aminoacid, aminoacyl, ONH2, O-aminoalkyl, O-aminoacid, O-aminoacyl, heterocycloalkyl, heterocycloalkaryl, aminoalkylamino, polyalklylamino, substituted silyl, or group having Formula I; R9 is O, S, CH2, S=O, CHF, or CF2.

In one embodiment, a siNA molecule of the invention comprises at least one (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) inverted abasic moiety, for example a compound having Formula VI:



wherein each R3, R4, R5, R6, R7, R8, R10, R11, R12, and R13 is independently H, OH, alkyl, substituted alkyl, alkaryl or aralkyl, F, Cl, Br, CN, CF3, OCF3, OCN, O-alkyl, S-alkyl, N-alkyl, O-alkyl, S-alkyl, N-alkyl, O-alkyl-OH, O-alkyl-OH, O-alkyl-SH, S-alkyl-OH, S-alkyl-SH, alkyl-S-alkyl, alkyl-O-alkyl, ONO2, NO2, N3, NH2, aminoalkyl, aminoacid, aminoacyl, ONH2, O-aminoalkyl, O-aminoacid, O-aminoacyl, heterocycloalkyl, heterocycloalkaryl, aminoalkylamino, polyalklylamino, substituted silyl, or group having Formula I; R9 is O, S, CH2, S=O, CHF, or CF2, and

5

10

either R2, R3, R8 or R13 serve as points of attachment to the siNA molecule of the invention.

In another embodiment, a siNA molecule of the invention comprises at least one (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) substituted polyalkyl moieties, for example a compound having Formula VII:

5

10

15

20

25

BNSDOCID: <WO\_

03074654A2\_l\_>

$$R_1$$
 $n$ 
 $R_2$ 

wherein each n is independently an integer from 1 to 12, each R1, R2 and R3 is independently H, OH, alkyl, substituted alkyl, alkaryl or aralkyl, F, Cl, Br, CN, CF3, OCF3, OCN, O-alkyl, S-alkyl, N-alkyl, O-alkenyl, S-alkenyl, N-alkenyl, SO-alkyl, alkyl-OSH, alkyl-OH, O-alkyl-OH, O-alkyl-SH, S-alkyl-OH, S-alkyl-SH, alkyl-S-alkyl, alkyl-O-alkyl, ONO2, NO2, N3, NH2, aminoalkyl, aminoacid, aminoacyl, ONH2, O-aminoacyl, O-aminoacyl, heterocycloalkyl, heterocycloalkaryl, aminoalkylamino, polyalklylamino, substituted silyl, or a group having Formula I, and R1, R2 or R3 serves as points of attachment to the siNA molecule of the invention.

In another embodiment, the invention features a compound having Formula VII, wherein R1 and R2 are hydroxyl (OH) groups, n=1, and R3 comprises O and is the point of attachment to the 3'-end, the 5'-end, or both of the 3' and 5'-ends of one or both strands of a double-stranded siNA molecule of the invention or to a single-stranded siNA molecule of the invention. This modification is referred to herein as "glyceryl" (for example modification 6 in **Figure 22**).

In another embodiment, a moiety having any of Formula V, VI or VII of the invention is at the 3'-end, the 5'-end, or both of the 3' and 5'-ends of a siNA molecule of the invention. For example, a moiety having Formula V, VI or VII can be present at the 3'-end, the 5'-end, or both of the 3' and 5'-ends of the antisense strand, the sense strand, or both antisense and sense strands of the siNA molecule. In addition, a moiety having Formula VII can be present at the 3'-end or the 5'-end of a hairpin siNA molecule as described herein.

In another embodiment, a siNA molecule of the invention comprises an abasic residue having Formula V or VI, wherein the abasic residue having Formula VI or VI is connected to the siNA construct in a 3'-3', 3'-2', 2'-3', or 5'-5' configuration, such as at the 3'-end, the 5'-end, or both of the 3' and 5'-ends of one or both siNA strands.

In one embodiment, a siNA molecule of the invention comprises one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) locked nucleic acid (LNA) nucleotides, for example at the 5'-end, the 3'-end, both of the 5' and 3'-ends, or any combination thereof, of the siNA molecule.

In another embodiment, a siNA molecule of the invention comprises one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) acyclic nucleotides, for example at the 5'-end, the 3'-end, both of the 5' and 3'-ends, or any combination thereof, of the siNA molecule.

In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule of the invention, wherein the chemically-modified siNA comprises a sense region, where any (e.g., one or more or all) pyrimidine nucleotides present in the sense region are 2'-deoxy-2'-fluoro pyrimidine nucleotides (e.g., wherein all pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides or alternately a plurality of pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides), and where any (e.g., one or more or all) purine nucleotides present in the sense region are 2'-deoxy purine nucleotides (e.g., wherein all purine nucleotides are 2'-deoxy purine nucleotides or alternately a plurality of purine nucleotides are 2'-deoxy purine nucleotides).

In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule of the invention, wherein the chemically-modified siNA comprises a sense region, where any (e.g., one or more or all) pyrimidine nucleotides present in the sense region are 2'-deoxy-2'-fluoro pyrimidine nucleotides (e.g., wherein all pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides or alternately a plurality of pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides), and where any (e.g., one or more or all) purine nucleotides present in the sense region are 2'-deoxy purine nucleotides (e.g., wherein all purine nucleotides are 2'-deoxy purine nucleotides or alternately a plurality of purine nucleotides are 2'-deoxy purine

5

10

15

20

25

nucleotides), wherein any nucleotides comprising a 3'-terminal nucleotide overhang that are present in said sense region are 2'-deoxy nucleotides.

In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule of the invention, wherein the chemically-modified siNA comprises an antisense region, where any (e.g., one or more or all) pyrimidine nucleotides present in the antisense region are 2'-deoxy-2'-fluoro pyrimidine nucleotides (e.g., wherein all pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides or alternately a plurality of pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides), and wherein any (e.g., one or more or all) purine nucleotides present in the antisense region are 2'-O-methyl purine nucleotides (e.g., wherein all purine nucleotides are 2'-O-methyl purine nucleotides or alternately a plurality of purine nucleotides are 2'-O-methyl purine nucleotides).

In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule of the invention, wherein the chemically-modified siNA comprises an antisense region, where any (e.g., one or more or all) pyrimidine nucleotides present in the antisense region are 2'-deoxy-2'-fluoro pyrimidine nucleotides (e.g., wherein all pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides or alternately a plurality of pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides), and wherein any (e.g., one or more or all) purine nucleotides present in the antisense region are 2'-O-methyl purine nucleotides (e.g., wherein all purine nucleotides are 2'-O-methyl purine nucleotides or alternately a plurality of purine nucleotides are 2'-O-methyl purine nucleotides), wherein any nucleotides comprising a 3'-terminal nucleotide overhang that are present in said antisense region are 2'-deoxy nucleotides.

In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule of the invention, wherein the chemically-modified siNA comprises an antisense region, where any (e.g., one or more or all) pyrimidine nucleotides present in the antisense region are 2'-deoxy-2'-fluoro pyrimidine nucleotides (e.g., wherein all pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides or alternately a plurality of pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides), and where any (e.g., one or more or all) purine nucleotides present in the antisense region are 2'-deoxy purine nucleotides (e.g., wherein all purine nucleotides are

5

10

15

20

25

2'-deoxy purine nucleotides or alternately a plurality of purine nucleotides are 2'-deoxy purine nucleotides).

In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule of the invention capable of mediating RNA interference (RNAi) inside a cell or reconstituted in vitro system, wherein the chemically-modified siNA comprises a sense region and an antisense region. The sense region comprises one 2'-deoxy-2'-fluoro pyrimidine nucleotides (e.g., wherein all pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides or alternately a plurality of pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides), and one or more 2'-deoxy purine nucleotides (e.g., wherein all purine nucleotides are 2'-deoxy purine nucleotides or alternately a plurality of purine nucleotides are 2'-deoxy purine nucleotides). Inverted deoxy abasic modifications can be optionally present at the 3'end, the 5'-end, or both of the 3' and 5'-ends of the sense region. The sense region optionally further comprises a 3'-terminal overhang having about 1 to about 4 (e.g., about 1, 2, 3, or 4) 2'-deoxyribonucleotides. The antisense region comprises one or more 2'deoxy-2'-fluoro pyrimidine nucleotides (e.g., wherein all pyrimidine nucleotides are 2'deoxy-2'-fluoro pyrimidine nucleotides or alternately a plurality of pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides), and one or more 2'-O-methyl purine nucleotides (e.g., wherein all purine nucleotides are 2'-O-methyl purine nucleotides or alternately a plurality of purine nucleotides are 2'-O-methyl purine nucleotides). A terminal cap modification, such as any modification described herein or shown in Figure 22, is optionally present at the 3'-end, the 5'-end, or both of the 3' and 5'-ends of the antisense sequence. The antisense region optionally further comprises a 3'-terminal nucleotide overhang having about 1 to about 4 (e.g., about 1, 2, 3, or 4) 2'deoxynucleotides, wherein the overhang nucleotides can further comprise one or more (e.g., 1, 2, 3, or 4) phosphorothioate internucleotide linkages. Non-limiting examples of these chemically-modified siNAs are shown in Figures 18 and 19 and Table IV herein.

In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule of the invention capable of mediating RNA interference (RNAi) inside a cell or reconstituted *in vitro* system, wherein the siNA comprises a sense region and an antisense region, wherein the sense region comprises one or more 2'-deoxy-2'-fluoro pyrimidine nucleotides (e.g., wherein all pyrimidine nucleotides are 2'-deoxy-2'-

5

10

15

20

25

fluoro pyrimidine nucleotides or alternately a plurality of pyrimidine nucleotides are 2'deoxy-2'-fluoro pyrimidine nucleotides), and one or more purine ribonucleotides (e.g., wherein all purine nucleotides are purine ribonucleotides or alternately a plurality of purine nucleotides are purine ribonucleotides) and wherein the antisense region comprises one or more 2'-deoxy-2'-fluoro pyrimidine nucleotides (e.g., wherein all pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides or alternately a plurality of pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides), and one or more 2'-O-methyl purine nucleotides (e.g., wherein all purine nucleotides are 2'-O-methyl purine nucleotides or alternately a plurality of purine nucleotides are 2'-O-methyl purine nucleotides). Inverted deoxy abasic modifications are optionally present at the 3'-end, the 5'-end, or both of the 3' and 5'-ends of the sense region. The sense region optionally further comprises a 3'-terminal overhang having about 1 to about 4 (e.g., about 1, 2, 3, or 4) 2'-deoxyribonucleotides. A terminal cap modification, such as any modification described herein or shown in Figure 22, is optionally present at the 3'-end, the 5'-end, or both of the 3' and 5'-ends of the antisense sequence. The antisense region optionally further comprises a 3'-terminal nucleotide overhang having about 1 to about 4 (e.g., about 1, 2, 3, or 4) 2'-deoxynucleotides, wherein the overhang nucleotides can further comprise one or more (e.g., 1, 2, 3, or 4) phosphorothioate internucleotide linkages. Non-limiting examples of these chemically-modified siNAs are shown in Figures 18 and 19 and Table IV herein.

In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule of the invention capable of mediating RNA interference (RNAi) inside a cell or reconstituted *in vitro* system, wherein the chemically-modified siNA comprises a sense region and an antisense region, wherein the sense region comprises one or 2'-deoxy-2'-fluoro pyrimidine nucleotides (e.g., wherein all pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides or alternately a plurality of pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides), and one or more purine nucleotides selected from the group consisting of 2'-deoxy nucleotides, locked nucleic acid (LNA) nucleotides, 2'-methoxyethyl nucleotides are selected from the group consisting of 2'-deoxy nucleotides, locked nucleic acid (LNA) nucleotides, 2'-methoxyethyl nucleotides, 4'-thionucleotides, 2'-methoxyethyl nucleotides, 4'-thionucleotides, 2'-methoxyethyl nucleotides, 4'-thionucleotides, and 2'-O-methyl nucleotides or alternately

5

10

15

20

25

a plurality of purine nucleotides are selected from the group consisting of 2'-deoxy nucleotides, locked nucleic acid (LNA) nucleotides, 2'-methoxyethyl nucleotides, 4'thionucleotides, and 2'-O-methyl nucleotides) and wherein the antisense region comprises one or more 2'-deoxy-2'-fluoro pyrimidine nucleotides (e.g., wherein all pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides or alternately a plurality of pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides), and one or more purine nucleotides selected from the group consisting of 2'-deoxy nucleotides, locked nucleic acid (LNA) nucleotides, 2'-methoxyethyl nucleotides, 4'-thionucleotides, and 2'-O-methyl nucleotides (e.g., wherein all purine nucleotides are selected from the group consisting of 2'-deoxy nucleotides, locked nucleic acid (LNA) nucleotides, 2'methoxyethyl nucleotides, 4'-thionucleotides, and 2'-O-methyl nucleotides or alternately a plurality of purine nucleotides are selected from the group consisting of 2'-deoxy nucleotides, locked nucleic acid (LNA) nucleotides, 2'-methoxyethyl nucleotides, 4'thionucleotides, and 2'-O-methyl nucleotides). Inverted deoxy abasic modifications are optionally present at the 3'-end, the 5'-end, or both of the 3' and 5'-ends of the sense region. The sense region optionally further comprises a 3'-terminal overhang having about 1 to about 4 (e.g., about 1, 2, 3, or 4) 2'-deoxyribonucleotides. A terminal cap modification, such as any modification described herein or shown in Figure 22, is optionally present at the 3'-end, the 5'-end, or both of the 3' and 5'-ends of the antisense sequence. The antisense region optionally further comprises a 3'-terminal nucleotide overhang having about 1 to about 4 (e.g., about 1, 2, 3, or 4) 2'-deoxynucleotides, wherein the overhang nucleotides can further comprise one or more (e.g., 1, 2, 3, or 4) phosphorothioate internucleotide linkages.

In another embodiment, any modified nucleotides present in the siNA molecules of the invention, preferably in the antisense strand of the siNA molecules of the invention, but also optionally in the sense and/or both antisense and sense strands, comprise modified nucleotides having properties or characteristics similar to naturally occurring ribonucleotides. For example, the invention features siNA molecules including modified nucleotides having a Northern conformation (e.g., Northern pseudorotation cycle, see for example Saenger, *Principles of Nucleic Acid Structure*, Springer-Verlag ed., 1984). As such, chemically modified nucleotides present in the siNA molecules of the invention, preferably in the antisense strand of the siNA molecules of the invention, but also

5

10

15

20

25

optionally in the sense and/or both antisense and sense strands, are resistant to nuclease degradation while at the same time maintaining the capacity to mediate RNAi. Non-limiting examples of nucleotides having a northern configuration include locked nucleic acid (LNA) nucleotides (e.g., 2'-O,4'-C-methylene-(D-ribofuranosyl) nucleotides); 2'-methoxyethoxy (MOE) nucleotides; 2'-methyl-thio-ethyl, 2'-deoxy-2'-fluoro nucleotides, 2'-deoxy-2'-chloro nucleotides, 2'-azido nucleotides, and 2'-O-methyl nucleotides.

In one embodiment, the invention features a chemically-modified short interfering nucleic acid molecule (siNA) capable of mediating RNA interference (RNAi) inside a cell or reconstituted in vitro system, wherein the chemical modification comprises a conjugate covalently attached to the chemically-modified siNA molecule. In another embodiment, the conjugate is covalently attached to the chemically-modified siNA molecule via a biodegradable linker. In one embodiment, the conjugate molecule is attached at the 3'end of either the sense strand, the antisense strand, or both strands of the chemicallymodified siNA molecule. In another embodiment, the conjugate molecule is attached at the 5'-end of either the sense strand, the antisense strand, or both strands of the chemically-modified siNA molecule. In yet another embodiment, the conjugate molecule is attached both the 3'-end and 5'-end of either the sense strand, the antisense strand, or both strands of the chemically-modified siNA molecule, or any combination thereof. In one embodiment, a conjugate molecule of the invention comprises a molecule that facilitates delivery of a chemically-modified siNA molecule into a biological system, In another embodiment, the conjugate molecule attached to the such as a cell. chemically-modified siNA molecule is a poly ethylene glycol, human serum albumin, or a ligand for a cellular receptor that can mediate cellular uptake. Examples of specific conjugate molecules contemplated by the instant invention that can be attached to chemically-modified siNA molecules are described in Vargeese et al., U.S. Serial No. 10/201,394, incorporated by reference herein. The type of conjugates used and the extent of conjugation of siNA molecules of the invention can be evaluated for improved pharmacokinetic profiles, bioavailability, and/or stability of siNA constructs while at the same time maintaining the ability of the siNA to mediate RNAi activity. As such, one skilled in the art can screen siNA constructs that are modified with various conjugates to determine whether the siNA conjugate complex possesses improved properties while

5

10

15

20

25

maintaining the ability to mediate RNAi, for example in animal models as are generally known in the art.

In one embodiment, the invention features a short interfering nucleic acid (siNA) molecule of the invention, wherein the siNA further comprises a nucleotide, nonnucleotide, or mixed nucleotide/non-nucleotide linker that joins the sense region of the siNA to the antisense region of the siNA. In one embodiment, a nucleotide linker of the invention can be a linker of  $\geq 2$  nucleotides in length, for example 3, 4, 5, 6, 7, 8, 9, or  $10^{\circ}$ nucleotides in length. In another embodiment, the nucleotide linker can be a nucleic acid aptamer. By "aptamer" or "nucleic acid aptamer" as used herein is meant a nucleic acid molecule that binds specifically to a target molecule wherein the nucleic acid molecule has sequence that comprises a sequence recognized by the target molecule in its natural setting. Alternately, an aptamer can be a nucleic acid molecule that binds to a target molecule where the target molecule does not naturally bind to a nucleic acid. The target molecule can be any molecule of interest. For example, the aptamer can be used to bind to a ligand-binding domain of a protein, thereby preventing interaction of the naturally occurring ligand with the protein. This is a non-limiting example and those in the art will recognize that other embodiments can be readily generated using techniques generally known in the art. (See, for example, Gold et al., 1995, Annu. Rev. Biochem., 64, 763; Brody and Gold, 2000, J. Biotechnol., 74, 5; Sun, 2000, Curr. Opin. Mol. Ther., 2, 100; Kusser, 2000, J. Biotechnol., 74, 27; Hermann and Patel, 2000, Science, 287, 820; and Jayasena, 1999, Clinical Chemistry, 45, 1628.)

In yet another embodiment, a non-nucleotide linker of the invention comprises abasic nucleotide, polyether, polyamine, polyamide, peptide, carbohydrate, lipid, polyhydrocarbon, or other polymeric compounds (e.g. polyethylene glycols such as those having between 2 and 100 ethylene glycol units). Specific examples include those described by Seela and Kaiser, Nucleic Acids Res. 1990, 18:6353 and Nucleic Acids Res. 1987, 15:3113; Cload and Schepartz, J. Am. Chem. Soc. 1991, 113:6324; Richardson and Schepartz, J. Am. Chem. Soc. 1991, 113:5109; Ma et al., Nucleic Acids Res. 1993, 21:2585 and Biochemistry 1993, 32:1751; Durand et al., Nucleic Acids Res. 1990, 18:6353; McCurdy et al., Nucleosides & Nucleotides 1991, 10:287; Jschke et al., Tetrahedron Lett. 1993, 34:301; Ono et al., Biochemistry 1991, 30:9914; Arnold et al., International Publication No. WO 89/02439; Usman et al., International Publication No.

5

10

15

20

25

WO 95/06731; Dudycz et al., International Publication No. WO 95/11910 and Ferentz and Verdine, J. Am. Chem. Soc. 1991, 113:4000, all hereby incorporated by reference herein. A "non-nucleotide" further means any group or compound that can be incorporated into a nucleic acid chain in the place of one or more nucleotide units, including either sugar and/or phosphate substitutions, and allows the remaining bases to exhibit their enzymatic activity. The group or compound can be abasic in that it does not contain a commonly recognized nucleotide base, such as adenosine, guanine, cytosine, uracil or thymine, for example at the C1 position of the sugar.

In one embodiment, the invention features a short interfering nucleic acid (siNA) molecule capable of mediating RNA interference (RNAi) inside a cell or reconstituted in vitro system, wherein one or both strands of the siNA molecule that are assembled from two separate oligonucleotides do not comprise any ribonucleotides. For example, a siNA molecule can be assembled from a single oligonculeotide where the sense and antisense regions of the siNA comprise separate oligonucleotides not having any ribonucleotides (e.g., nucleotides having a 2'-OH group) present in the oligonucleotides. In another example, a siNA molecule can be assembled from a single oligonculeotide where the sense and antisense regions of the siNA are linked or circularized by a nucleotide or nonnucleotide linker as desrcibed herein, wherein the oligonucleotide does not have any ribonucleotides (e.g., nucleotides having a 2'-OH group) present in the oligonucleotide. Applicant has surprisingly found that the presense of ribonucleotides (e.g., nucleotides having a 2'-hydroxyl group) within the siNA molecule is not required or essential to support RNAi activity. As such, in one embodiment, all positions within the siNA can include chemically modified nucleotides and/or non-nucleotides such as nucleotides and or non-nucleotides having Formula I, II, III, IV, V, VI, or VII or any combination thereof to the extent that the ability of the siNA molecule to support RNAi activity in a cell is maintained.

In one embodiment, a siNA molecule of the invention is a single stranded siNA molecule that mediates RNAi activity in a cell or reconstituted in vitro system, wherein the siNA molecule comprises a single stranded polynucleotide having complementarity to a target nucleic acid sequence. In another embodiment, the single stranded siNA molecule of the invention comprises a 5'-terminal phosphate group. In another embodiment, the single stranded siNA molecule of the invention comprises a 5'-terminal

5

10

15

20

25

phosphate group and a 3'-terminal phosphate group (e.g., a 2', 3'-cyclic phosphate). In another embodiment, the single stranded siNA molecule of the invention comprises about 19 to about 29 nucleotides. In yet another embodiment, the single stranded siNA molecule of the invention comprises one or more chemically modified nucleotides or non-nucleotides described herein. For example, all the positions within the siNA molecule can include chemically-modified nucleotides such as nucleotides having any of Formulae I-VII, or any combination thereof to the extent that the ability of the siNA molecule to support RNAi activity in a cell is maintained.

In one embodiment, a siNA molecule of the invention is a single stranded siNA molecule that mediates RNAi activity in a cell or reconstituted in vitro system, wherein the siNA molecule comprises a single stranded polynucleotide having complementarity to a target nucleic acid sequence, and wherein one or more pyrimidine nucleotides present in the siNA are 2'-deoxy-2'-fluoro pyrimidine nucleotides (e.g., wherein all pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides or alternately a plurality of pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides), and wherein any purine nucleotides present in the antisense region are 2'-O-methyl purine nucleotides (e.g., wherein all purine nucleotides are 2'-O-methyl purine nucleotides or alternately a plurality of purine nucleotides are 2'-O-methyl purine nucleotides), and a terminal cap modification, such as any modification described herein or shown in Figure 22, that is optionally present at the 3'-end, the 5'-end, or both of the 3' and 5'-ends of the antisense sequence, the siNA optionally further comprising about 1 to about 4 (e.g., about 1, 2, 3, or 4) terminal 2'-deoxynucleotides at the 3'-end of the siNA molecule, wherein the terminal nucleotides can further comprise one or more (e.g., 1, 2, 3, or 4) phosphorothioate internucleotide linkages, and wherein the siNA optionally further comprises a terminal phosphate group, such as a 5'-terminal phosphate group.

In one embodiment, a siNA molecule of the invention is a single stranded siNA molecule that mediates RNAi activity in a cell or reconstituted in vitro system, wherein the siNA molecule comprises a single stranded polynucleotide having complementarity to a target nucleic acid sequence, and wherein one or more pyrimidine nucleotides present in the siNA are 2'-deoxy-2'-fluoro pyrimidine nucleotides (e.g., wherein all pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides or alternately a plurality of pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides), and wherein any

5

10

15

20

25

purine nucleotides present in the antisense region are 2'-deoxy purine nucleotides (e.g., wherein all purine nucleotides are 2'-deoxy purine nucleotides or alternately a plurality of purine nucleotides are 2'-deoxy purine nucleotides), and a terminal cap modification, such as any modification described herein or shown in **Figure 22**, that is optionally present at the 3'-end, the 5'-end, or both of the 3' and 5'-ends of the antisense sequence, the siNA optionally further comprising about 1 to about 4 (e.g., about 1, 2, 3, or 4) terminal 2'-deoxynucleotides at the 3'-end of the siNA molecule, wherein the terminal nucleotides can further comprise one or more (e.g., 1, 2, 3, or 4) phosphorothioate internucleotide linkages, and wherein the siNA optionally further comprises a terminal phosphate group, such as a 5'-terminal phosphate group.

In one embodiment, a siNA molecule of the invention is a single stranded siNA molecule that mediates RNAi activity in a cell or reconstituted in vitro system, wherein the siNA molecule comprises a single stranded polynucleotide having complementarity to a target nucleic acid sequence, and wherein one or more pyrimidine nucleotides present in the siNA are 2'-deoxy-2'-fluoro pyrimidine nucleotides (e.g., wherein all pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides or alternately a plurality of pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides), and wherein any purine nucleotides present in the antisense region are locked nucleic acid (LNA) nucleotides (e.g., wherein all purine nucleotides are LNA nucleotides or alternately a plurality of purine nucleotides are LNA nucleotides), and a terminal cap modification, such as any modification described herein or shown in Figure 22, that is optionally present at the 3'-end, the 5'-end, or both of the 3' and 5'-ends of the antisense sequence, the siNA optionally further comprising about 1 to about 4 (e.g., about 1, 2, 3, or 4) terminal 2'-deoxynucleotides at the 3'-end of the siNA molecule, wherein the terminal nucleotides can further comprise one or more (e.g., 1, 2, 3, or 4 ) phosphorothioate internucleotide linkages, and wherein the siNA optionally further comprises a terminal phosphate group, such as a 5'-terminal phosphate group.

In one embodiment, a siNA molecule of the invention is a single stranded siNA molecule that mediates RNAi activity in a cell or reconstituted in vitro system, wherein the siNA molecule comprises a single stranded polynucleotide having complementarity to a target nucleic acid sequence, and wherein one or more pyrimidine nucleotides present in the siNA are 2'-deoxy-2'-fluoro pyrimidine nucleotides (e.g., wherein all pyrimidine

5

10

15

20

25

nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides or alternately a plurality of pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides), and wherein any purine nucleotides present in the antisense region are 2'-methoxyethyl purine nucleotides (e.g., wherein all purine nucleotides are 2'-methoxyethyl purine nucleotides or alternately a plurality of purine nucleotides are 2'-methoxyethyl purine nucleotides), and a terminal cap modification, such as any modification described herein or shown in **Figure 22**, that is optionally present at the 3'-end, the 5'-end, or both of the 3' and 5'-ends of the antisense sequence, the siNA optionally further comprising about 1 to about 4 (e.g., about 1, 2, 3, or 4) terminal 2'-deoxynucleotides at the 3'-end of the siNA molecule, wherein the terminal nucleotides can further comprise one or more (e.g., 1, 2, 3, or 4) phosphorothioate internucleotide linkages, and wherein the siNA optionally further comprises a terminal phosphate group, such as a 5'-terminal phosphate group.

In another embodiment, any modified nucleotides present in the single stranded siNA molecules of the invention comprise modified nucleotides having properties or characteristics similar to naturally occurring ribonucleotides. For example, the invention features siNA molecules including modified nucleotides having a Northern conformation (e.g., Northern pseudorotation cycle, see for example Saenger, *Principles of Nucleic Acid Structure*, Springer-Verlag ed., 1984). As such, chemically modified nucleotides present in the single stranded siNA molecules of the invention are preferably resistant to nuclease degradation while at the same time maintaining the capacity to mediate RNAi.

In one embodiment, the invention features a method for modulating the expression of a gene within a cell comprising: (a) synthesizing a siNA molecule of the invention, which can be chemically-modified, wherein one of the siNA strands comprises a sequence complementary to RNA of the gene; and (b) introducing the siNA molecule into a cell under conditions suitable to modulate the expression of the gene in the cell.

In one embodiment, the invention features a method for modulating the expression of a gene within a cell comprising: (a) synthesizing a siNA molecule of the invention, which can be chemically-modified, wherein one of the siNA strands comprises a sequence complementary to RNA of the gene and wherein the sense strand sequence of the siNA comprises a sequence substantially similar to the sequence of the target RNA;

5

10

15

20

25

and (b) introducing the siNA molecule into a cell under conditions suitable to modulate the expression of the gene in the cell.

In another embodiment, the invention features a method for modulating the expression of more than one gene within a cell comprising: (a) synthesizing siNA molecules of the invention, which can be chemically-modified, wherein one of the siNA strands comprises a sequence complementary to RNA of the genes; and (b) introducing the siNA molecules into a cell under conditions suitable to modulate the expression of the genes in the cell.

In another embodiment, the invention features a method for modulating the expression of more than one gene within a cell comprising: (a) synthesizing a siNA molecule of the invention, which can be chemically-modified, wherein one of the siNA strands comprises a sequence complementary to RNA of the gene and wherein the sense strand sequence of the siNA comprises a sequence substantially similar to the sequence of the target RNA; and (b) introducing the siNA molecules into a cell under conditions suitable to modulate the expression of the genes in the cell.

In one embodiment, siNA molecules of the invention are used as reagents in ex vivo applications. For example, siNA reagents are intoduced into tissue or cells that are transplanted into a subject for therapeutic effect. The cells and/or tissue can be derived from an organism or subject that later receives the explant, or can be derived from another organism or subject prior to transplantation. The siNA molecules can be used to modulate the expression of one or more genes in the cells or tissue, such that the cells or tissue obtain a desired phenotype or are able to perform a function when transplanted in In one embodiment, certain target cells from a patient are extracted. extracted cells are contacted with siNAs targeteing a specific nucleotide sequence within the cells under conditions suitable for uptake of the siNAs by these cells (e.g. using delivery reagents such as cationic lipids, liposomes and the like or using techniques such as electroporation to facilitate the delivery of siNAs into cells). The cells are then reintroduced back into the same patient or other patients. Non-limiting examples of ex vivo applications include use in organ/tissue transplant, tissue grafting, or treatment of pulmonary disease (e.g., restenosis) or prevent neointimal hyperplasia and atherosclerosis in vein grafts. Such ex vivo applications may also used to treat conditions associated with

5

10

15

20

25

coronary and peripheral bypass graft failure, for example, such methods can be used in conjunction with peripheral vascular bypass graft surgery and coronary artery bypass graft surgery. Additional applications include transplants to treat CNS lesions or injury, including use in treatment of neurodegenerative conditions such as Alzheimer's disease, Parkinson's Disease, Epilepsy, Dementia, Huntington's disease, or amyotrophic lateral sclerosis (ALS).

In one embodiment, the invention features a method of modulating the expression of a gene in a tissue explant comprising: (a) synthesizing a siNA molecule of the invention, which can be chemically-modified, wherein one of the siNA strands comprises a sequence complementary to RNA of the gene; and (b) introducing the siNA molecule into a cell of the tissue explant derived from a particular organism under conditions suitable to modulate the expression of the gene in the tissue explant. In another embodiment, the method further comprises introducing the tissue explant back into the organism the tissue was derived from or into another organism under conditions suitable to modulate the expression of the gene in that organism.

In one embodiment, the invention features a method of modulating the expression of a gene in a tissue explant comprising: (a) synthesizing a siNA molecule of the invention, which can be chemically-modified, wherein one of the siNA strands comprises a sequence complementary to RNA of the gene and wherein the sense strand sequence of the siNA comprises a sequence substantially similar to the sequence of the target RNA; and (b) introducing the siNA molecule into a cell of the tissue explant derived from a particular organism under conditions suitable to modulate the expression of the gene in the tissue explant. In another embodiment, the method further comprises introducing the tissue explant back into the organism the tissue was derived from or into another organism under conditions suitable to modulate the expression of the gene in that organism.

In another embodiment, the invention features a method of modulating the expression of more than one gene in a tissue explant comprising: (a) synthesizing siNA molecules of the invention, which can be chemically-modified, wherein one of the siNA strands comprises a sequence complementary to RNA of the genes; and (b) introducing the siNA molecules into a cell of the tissue explant derived from a particular organism

5

10

15

20

25

under conditions suitable to modulate the expression of the genes in the tissue explant. In another embodiment, the method further comprises introducing the tissue explant back into the organism the tissue was derived from or into another organism under conditions suitable to modulate the expression of the genes in that organism.

In one embodiment, the invention features a method of modulating the expression of a gene in an organism comprising: (a) synthesizing a siNA molecule of the invention, which can be chemically-modified, wherein one of the siNA strands comprises a sequence complementary to RNA of the gene; and (b) introducing the siNA molecule into the organism under conditions suitable to modulate the expression of the gene in the organism.

In another embodiment, the invention features a method of modulating the expression of more than one gene in an organism comprising: (a) synthesizing siNA molecules of the invention, which can be chemically-modified, wherein one of the siNA strands comprises a sequence complementary to RNA of the genes; and (b) introducing the siNA molecules into the organism under conditions suitable to modulate the expression of the genes in the organism.

In one embodiment, the invention features a method for modulating the expression of a gene within a cell comprising: (a) synthesizing a siNA molecule of the invention, which can be chemically-modified, wherein the siNA comprises a single stranded sequence having complementarity to RNA of the gene; and (b) introducing the siNA molecule into a cell under conditions suitable to modulate the expression of the gene in the cell.

In another embodiment, the invention features a method for modulating the expression of more than one gene within a cell comprising: (a) synthesizing siNA molecules of the invention, which can be chemically-modified, wherein the siNA comprises a single stranded sequence having complementarity to RNA of the gene; and (b) contacting the siNA molecule with a cell in vitro or in vivo under conditions suitable to modulate the expression of the genes in the cell.

In one embodiment, the invention features a method of modulating the expression of a gene in a tissue explant comprising: (a) synthesizing a siNA molecule of the

5

10

15

20

25

invention, which can be chemically-modified, wherein the siNA comprises a single stranded sequence having complementarity to RNA of the gene; and (b) contacting the siNA molecule with a cell of the tissue explant derived from a particular organism under conditions suitable to modulate the expression of the gene in the tissue explant. In another embodiment, the method further comprises introducing the tissue explant back into the organism the tissue was derived from or into another organism under conditions suitable to modulate the expression of the gene in that organism.

In another embodiment, the invention features a method of modulating the expression of more than one gene in a tissue explant comprising: (a) synthesizing siNA molecules of the invention, which can be chemically-modified, wherein the siNA comprises a single stranded sequence having complementarity to RNA of the gene; and (b) introducing the siNA molecules into a cell of the tissue explant derived from a particular organism under conditions suitable to modulate the expression of the genes in the tissue explant. In another embodiment, the method further comprises introducing the tissue explant back into the organism the tissue was derived from or into another organism under conditions suitable to modulate the expression of the genes in that organism.

In one embodiment, the invention features a method of modulating the expression of a gene in an organism comprising: (a) synthesizing a siNA molecule of the invention, which can be chemically-modified, wherein the siNA comprises a single stranded sequence having complementarity to RNA of the gene; and (b) introducing the siNA molecule into the organism under conditions suitable to modulate the expression of the gene in the organism.

In another embodiment, the invention features a method of modulating the expression of more than one gene in an organism comprising: (a) synthesizing siNA molecules of the invention, which can be chemically-modified, wherein the siNA comprises a single stranded sequence having complementarity to RNA of the gene; and (b) introducing the siNA molecules into the organism under conditions suitable to modulate the expression of the genes in the organism.

In one embodiment, the invention features a method of modulating the expression of a gene in an organism comprising contacting the organism with a siNA molecule of the

5

10

15

20

25

invention under conditions suitable to modulate the expression of the gene in the organism.

In another embodiment, the invention features a method of modulating the expression of more than one gene in an organism comprising contacting the organism with one or more siNA molecules of the invention under conditions suitable to modulate the expression of the genes in the organism.

The siNA molecules of the invention can be designed to inhibit target gene expression through RNAi targeting of a variety of RNA molecules. In one embodiment, the siNA molecules of the invention are used to target various RNAs corresponding to a target gene. Non-limiting examples of such RNAs include messenger RNA (mRNA), alternate RNA splice variants of target gene(s), post-transcriptionally modified RNA of target gene(s), pre-mRNA of target gene(s), and/or RNA templates. If alternate splicing produces a family of transcripts that are distinguished by usage of appropriate exons, the instant invention can be used to inhibit gene expression through the appropriate exons to specifically inhibit or to distinguish among the functions of gene family members. For example, a protein that contains an alternatively spliced transmembrane domain can be expressed in both membrane bound and secreted forms. Use of the invention to target the exon containing the transmembrane domain can be used to determine the functional consequences of pharmaceutical targeting of membrane bound as opposed to the secreted form of the protein. Non-limiting examples of applications of the invention relating to targeting these RNA molecules include therapeutic pharmaceutical applications, pharmaceutical discovery applications, molecular diagnostic and gene function applications, and gene mapping, for example using single nucleotide polymorphism mapping with siNA molecules of the invention. Such applications can be implemented using known gene sequences or from partial sequences available from an expressed sequence tag (EST).

In another embodiment, the siNA molecules of the invention are used to target conserved sequences corresponding to a gene family or gene families. As such, siNA molecules targeting multiple gene targets can provide increased therapeutic effect. In addition, siNA can be used to characterize pathways of gene function in a variety of applications. For example, the present invention can be used to inhibit the activity of

5

10

15

20

25

target gene(s) in a pathway to determine the function of uncharacterized gene(s) in gene function analysis, mRNA function analysis, or translational analysis. The invention can be used to determine potential target gene pathways involved in various diseases and conditions toward pharmaceutical development. The invention can be used to understand pathways of gene expression involved in, for example, in development, such as prenatal development and postnatal development, and/or the progression and/or maintenance of cancer, infectious disease, autoimmunity, inflammation, endocrine disorders, renal disease, pulmonary disease, cardiovascular disease, birth defects, ageing, any other disease or condition related to gene expression.

In one embodiment, the invention features a method comprising: (a) generating a library of siNA constructs having a predetermined complexity; and (b) assaying the siNA constructs of (a) above, under conditions suitable to determine RNAi target sites within the target RNA sequence. In another embodiment, the siNA molecules of (a) have strands of a fixed length, for example, about 23 nucleotides in length. In yet another embodiment, the siNA molecules of (a) are of differing length, for example having strands of about 19 to about 25 (e.g., about 19, 20, 21, 22, 23, 24, or 25) nucleotides in length. In one embodiment, the assay can comprise a reconstituted *in vitro* siNA assay as described herein. In another embodiment, the assay can comprise a cell culture system in which target RNA is expressed. In another embodiment, fragments of target RNA are analyzed for detectable levels of cleavage, for example by gel electrophoresis, northern blot analysis, or RNAse protection assays, to determine the most suitable target site(s) within the target RNA sequence. The target RNA sequence can be obtained as is known in the art, for example, by cloning and/or transcription for *in vitro* systems, and by cellular expression in *in vivo* systems.

In one embodiment, the invention features a method comprising: (a) generating a randomized library of siNA constructs having a predetermined complexity, such as of 4<sup>N</sup>, where N represents the number of base paired nucleotides in each of the siNA construct strands (eg. for a siNA construct having 21 nucleotide sense and antisense strands with 19 base pairs, the complexity would be 4<sup>19</sup>); and (b) assaying the siNA constructs of (a) above, under conditions suitable to determine RNAi target sites within the target RNA sequence. In another embodiment, the siNA molecules of (a) have strands of a fixed length, for example about 23 nucleotides in length. In yet another embodiment, the siNA

5

10

15

20

25

molecules of (a) are of differing length, for example having strands of about 19 to about 25 (e.g., about 19, 20, 21, 22, 23, 24, or 25) nucleotides in length. In one embodiment, the assay can comprise a reconstituted *in vitro* siNA assay as described in Example 7 herein. In another embodiment, the assay can comprise a cell culture system in which target RNA is expressed. In another embodiment, fragments of target RNA are analyzed for detectable levels of cleavage, for example by gel electrophoresis, northern blot analysis, or RNAse protection assays, to determine the most suitable target site(s) within the target RNA sequence. In another embodiment, the target RNA sequence can be obtained as is known in the art, for example, by cloning and/or transcription for *in vitro* systems, and by cellular expression in *in vivo* systems.

In another embodiment, the invention features a method comprising: (a) analyzing the sequence of a RNA target encoded by a target gene; (b) synthesizing one or more sets of siNA molecules having sequence complementary to one or more regions of the RNA of (a); and (c) assaying the siNA molecules of (b) under conditions suitable to determine RNAi targets within the target RNA sequence. In one embodiment, the siNA molecules of (b) have strands of a fixed length, for example about 23 nucleotides in length. In another embodiment, the siNA molecules of (b) are of differing length, for example having strands of about 19 to about 25 (e.g., about 19, 20, 21, 22, 23, 24, or 25) nucleotides in length. In one embodiment, the assay can comprise a reconstituted in vitro siNA assay as described herein. In another embodiment, the assay can comprise a cell culture system in which target RNA is expressed. Fragments of target RNA are analyzed for detectable levels of cleavage, for example by gel electrophoresis, northern blot analysis, or RNAse protection assays, to determine the most suitable target site(s) within the target RNA sequence. The target RNA sequence can be obtained as is known in the art, for example, by cloning and/or transcription for in vitro systems, and by expression in in vivo systems.

By "target site" is meant a sequence within a target RNA that is "targeted" for cleavage mediated by a siNA construct which contains sequences within its antisense region that are complementary to the target sequence.

By "detectable level of cleavage" is meant cleavage of target RNA (and formation of cleaved product RNAs) to an extent sufficient to discern cleavage products above the

5

10

15

20

25

background of RNAs produced by random degradation of the target RNA. Production of cleavage products from 1-5% of the target RNA is sufficient to detect above the background for most methods of detection.

In one embodiment, the invention features a composition comprising a siNA molecule of the invention, which can be chemically-modified, in a pharmaceutically acceptable carrier or diluent. In another embodiment, the invention features a pharmaceutical composition comprising siNA molecules of the invention, which can be chemically-modified, targeting one or more genes in a pharmaceutically acceptable carrier or diluent. In another embodiment, the invention features a method for treating or preventing a disease or condition in a subject, comprising administering to the subject a composition of the invention under conditions suitable for the treatment or prevention of the disease or condition in the subject, alone or in conjunction with one or more other therapeutic compounds. In yet another embodiment, the invention features a method for reducing or preventing tissue rejection in a subject comprising administering to the subject a composition of the invention under conditions suitable for the reduction or prevention of tissue rejection in the subject.

In another embodiment, the invention features a method for validating a gene target, comprising: (a) synthesizing a siNA molecule of the invention, which can be chemically-modified, wherein one of the siNA strands includes a sequence complementary to RNA of a target gene; (b) introducing the siNA molecule into a cell, tissue, or organism under conditions suitable for modulating expression of the target gene in the cell, tissue, or organism; and (c) determining the function of the gene by assaying for any phenotypic change in the cell, tissue, or organism.

In another embodiment, the invention features a method for validating a target gene comprising: (a) synthesizing a siNA molecule of the invention, which can be chemically-modified, wherein one of the siNA strands includes a sequence complementary to RNA of a target gene; (b) introducing the siNA molecule into a biological system under conditions suitable for modulating expression of the target gene in the biological system; and (c) determining the function of the gene by assaying for any phenotypic change in the biological system.

5

10

15

20

25

By "biological system" is meant, material, in a purified or unpurified form, from biological sources, including but not limited to human, animal, plant, insect, bacterial, viral or other sources, wherein the system comprises the components required for RNAi acitivity. The term "biological system" includes, for example, a cell, tissue, or organism, or extract thereof. The term biological system also includes reconstituted RNAi systems that can be used in an *in vitro* setting.

5

10

15

20

25

30

03074654A2 L>

BNSDOCID: <WO

By "phenotypic change" is meant any detectable change to a cell that occurs in response to contact or treatment with a nucleic acid molecule of the invention (e.g., siNA). Such detectable changes include, but are not limited to, changes in shape, size, proliferation, motility, protein expression or RNA expression or other physical or chemical changes as can be assayed by methods known in the art. The detectable change can also include expression of reporter genes/molecules such as Green Florescent Protein (GFP) or various tags that are used to identify an expressed protein or any other cellular component that can be assayed.

In one embodiment, the invention features a kit containing a siNA molecule of the invention, which can be chemically-modified, that can be used to modulate the expression of a target gene in a cell, tissue, or organism. In another embodiment, the invention features a kit containing more than one siNA molecule of the invention, which can be chemically-modified, that can be used to modulate the expression of more than one target gene in a cell, tissue, or organism.

In one embodiment, the invention features a kit containing a siNA molecule of the invention, which can be chemically-modified, that can be used to modulate the expression of a target gene in a biological system. In another embodiment, the invention features a kit containing more than one siNA molecule of the invention, which can be chemically-modified, that can be used to modulate the expression of more than one target gene in a biological system.

In one embodiment, the invention features a cell containing one or more siNA molecules of the invention, which can be chemically-modified. In another embodiment, the cell containing a siNA molecule of the invention is a mammalian cell. In yet another embodiment, the cell containing a siNA molecule of the invention is a human cell.

In one embodiment, the synthesis of a siNA molecule of the invention, which can be chemically-modified, comprises: (a) synthesis of two complementary strands of the siNA molecule; (b) annealing the two complementary strands together under conditions suitable to obtain a double-stranded siNA molecule. In another embodiment, synthesis of the two complementary strands of the siNA molecule is by solid phase oligonucleotide synthesis. In yet another embodiment, synthesis of the two complementary strands of the siNA molecule is by solid phase tandem oligonucleotide synthesis.

In one embodiment, the invention features a method for synthesizing a siNA duplex molecule comprising: (a) synthesizing a first oligonucleotide sequence strand of the siNA molecule, wherein the first oligonucleotide sequence strand comprises a cleavable linker molecule that can be used as a scaffold for the synthesis of the second oligonucleotide sequence strand of the siNA; (b) synthesizing the second oligonucleotide sequence strand of siNA on the scaffold of the first oligonucleotide sequence strand, wherein the second oligonucleotide sequence strand further comprises a chemical moiety than can be used to purify the siNA duplex; (c) cleaving the linker molecule of (a) under conditions suitable for the two siNA oligonucleotide strands to hybridize and form a stable duplex; and (d) purifying the siNA duplex utilizing the chemical moiety of the second oligonucleotide sequence strand. In one embodiment, cleavage of the linker molecule in (c) above takes place during deprotection of the oligonucleotide, for example under hydrolysis conditions using an alkylamine base such as methylamine. In one embodiment, the method of synthesis comprises solid phase synthesis on a solid support such as controlled pore glass (CPG) or polystyrene, wherein the first sequence of (a) is synthesized on a cleavable linker, such as a succinyl linker, using the solid support as a scaffold. The cleavable linker in (a) used as a scaffold for synthesizing the second strand can comprise similar reactivity as the solid support derivatized linker, such that cleavage of the solid support derivatized linker and the cleavable linker of (a) takes place concomitantly. In another embodiment, the chemical moiety of (b) that can be used to isolate the attached oligonucleotide sequence comprises a trityl group, for example a dimethoxytrityl group, which can be employed in a trityl-on synthesis strategy as described herein. In yet another embodiment, the chemical moiety, such as a dimethoxytrityl group, is removed during purification, for example, using acidic conditions.

5

10

15

20

25

In a further embodiment, the method for siNA synthesis is a solution phase synthesis or hybrid phase synthesis wherein both strands of the siNA duplex are synthesized in tandem using a cleavable linker attached to the first sequence which acts a scaffold for synthesis of the second sequence. Cleavage of the linker under conditions suitable for hybridization of the separate siNA sequence strands results in formation of the double-stranded siNA molecule.

In another embodiment, the invention features a method for synthesizing a siNA duplex molecule comprising: (a) synthesizing one oligonucleotide sequence strand of the siNA molecule, wherein the sequence comprises a cleavable linker molecule that can be used as a scaffold for the synthesis of another oligonucleotide sequence; (b) synthesizing a second oligonucleotide sequence having complementarity to the first sequence strand on the scaffold of (a), wherein the second sequence comprises the other strand of the doublestranded siNA molecule and wherein the second sequence further comprises a chemical moiety than can be used to isolate the attached oligonucleotide sequence; (c) purifying the product of (b) utilizing the chemical moiety of the second oligonucleotide sequence strand under conditions suitable for isolating the full-length sequence comprising both siNA oligonucleotide strands connected by the cleavable linker and under conditions suitable for the two siNA oligonucleotide strands to hybridize and form a stable duplex. In one embodiment, cleavage of the linker molecule in (c) above takes place during deprotection of the oligonucleotide, for example under hydrolysis conditions. In another embodiment, cleavage of the linker molecule in (c) above takes place after deprotection of the oligonucleotide. In another embodiment, the method of synthesis comprises solid phase synthesis on a solid support such as controlled pore glass (CPG) or polystyrene, wherein the first sequence of (a) is synthesized on a cleavable linker, such as a succinyl linker, using the solid support as a scaffold. The cleavable linker in (a) used as a scaffold for synthesizing the second strand can comprise similar reactivity or differing reactivity as the solid support derivatized linker, such that cleavage of the solid support derivatized linker and the cleavable linker of (a) takes place either concomitantly or sequentially. In one embodiment, the chemical moiety of (b) that can be used to isolate the attached oligonucleotide sequence comprises a trityl group, for example a dimethoxytrityl group.

In another embodiment, the invention features a method for making a doublestranded siNA molecule in a single synthetic process comprising: (a) synthesizing an

5

10

15

20

25

oligonucleotide having a first and a second sequence, wherein the first sequence is complementary to the second sequence, and the first oligonucleotide sequence is linked to the second sequence via a cleavable linker, and wherein a terminal 5'-protecting group, for example, a 5'-O-dimethoxytrityl group (5'-O-DMT) remains on the oligonucleotide having the second sequence; (b) deprotecting the oligonucleotide whereby the deprotection results in the cleavage of the linker joining the two oligonucleotide sequences; and (c) purifying the product of (b) under conditions suitable for isolating the double-stranded siNA molecule, for example using a trityl-on synthesis strategy as described herein.

In another embodiment, the method of synthesis of siNA molecules of the invention comprises the teachings of Scaringe *et al.*, US Patent Nos. 5,889,136; 6,008,400; and 6,111,086, incorporated by reference herein in their entirety.

In one embodiment, the invention features siNA constructs that mediate RNAi in a cell or reconstituted system, wherein the siNA construct comprises one or more chemical modifications, for example, one or more chemical modifications having any of Formulae I-VII or any combination thereof that increases the nuclease resistance of the siNA construct.

In another embodiment, the invention features a method for generating siNA molecules with increased nuclease resistance comprising (a) introducing nucleotides having any of Formula I-VII or any combination thereof into a siNA molecule, and (b) assaying the siNA molecule of step (a) under conditions suitable for isolating siNA molecules having increased nuclease resistance.

In one embodiment, the invention features siNA constructs that mediate RNAi against a target gene, wherein the siNA construct comprises one or more chemical modifications described herein that modulates the binding affinity between the sense and antisense strands of the siNA construct.

In another embodiment, the invention features a method for generating siNA molecules with increased binding affinity between the sense and antisense strands of the siNA molecule comprising (a) introducing nucleotides having any of Formula I-VII or any combination thereof into a siNA molecule, and (b) assaying the siNA molecule of

5

10

15

20

25

step (a) under conditions suitable for isolating siNA molecules having increased binding affinity between the sense and antisense strands of the siNA molecule.

In one embodiment, the invention features siNA constructs that mediate RNAi in a cell or reconstituted system, wherein the siNA construct comprises one or more chemical modifications described herein that modulates the binding affinity between the antisense strand of the siNA construct and a complementary target RNA sequence within a cell.

In one embodiment, the invention features siNA constructs that mediate RNAi in a cell or reconstituted system, wherein the siNA construct comprises one or more chemical modifications described herein that modulates the binding affinity between the antisense strand of the siNA construct and a complementary target DNA sequence within a cell.

In another embodiment, the invention features a method for generating siNA molecules with increased binding affinity between the antisense strand of the siNA molecule and a complementary target RNA sequence comprising (a) introducing nucleotides having any of Formula I-VII or any combination thereof into a siNA molecule, and (b) assaying the siNA molecule of step (a) under conditions suitable for isolating siNA molecules having increased binding affinity between the antisense strand of the siNA molecule and a complementary target RNA sequence.

In another embodiment, the invention features a method for generating siNA molecules with increased binding affinity between the antisense strand of the siNA molecule and a complementary target DNA sequence comprising (a) introducing nucleotides having any of Formula I-VII or any combination thereof into a siNA molecule, and (b) assaying the siNA molecule of step (a) under conditions suitable for isolating siNA molecules having increased binding affinity between the antisense strand of the siNA molecule and a complementary target DNA sequence.

In one embodiment, the invention features siNA constructs that mediate RNAi in a cell or reconstituted system, wherein the siNA construct comprises one or more chemical modifications described herein that modulate the polymerase activity of a cellular polymerase capable of generating additional endogenous siNA molecules having sequence homology to the chemically-modified siNA construct.

5

10

15

20

In another embodiment, the invention features a method for generating siNA molecules capable of mediating increased polymerase activity of a cellular polymerase capable of generating additional endogenous siNA molecules having sequence homology to a chemically-modified siNA molecule comprising (a) introducing nucleotides having any of Formula I-VII or any combination thereof into a siNA molecule, and (b) assaying the siNA molecule of step (a) under conditions suitable for isolating siNA molecules capable of mediating increased polymerase activity of a cellular polymerase capable of generating additional endogenous siNA molecules having sequence homology to the chemically-modified siNA molecule.

In one embodiment, the invention features chemically-modified siNA constructs that mediate RNAi in a cell or reconstituted system, wherein the chemical modifications do not significantly effect the interaction of siNA with a target RNA molecule, DNA molecule and/or proteins or other factors that are essential for RNAi in a manner that would decrease the efficacy of RNAi mediated by such siNA constructs.

In another embodiment, the invention features a method for generating siNA molecules with improved RNAi activity, comprising (a) introducing nucleotides having any of Formula I-VII or any combination thereof into a siNA molecule, and (b) assaying the siNA molecule of step (a) under conditions suitable for isolating siNA molecules having improved RNAi activity.

In yet another embodiment, the invention features a method for generating siNA molecules with improved RNAi activity against a target RNA comprising (a) introducing nucleotides having any of Formula I-VII or any combination thereof into a siNA molecule, and (b) assaying the siNA molecule of step (a) under conditions suitable for isolating siNA molecules having improved RNAi activity against the target RNA.

In yet another embodiment, the invention features a method for generating siNA molecules with improved RNAi activity against a DNA target comprising (a) introducing nucleotides having any of Formula I-VII or any combination thereof into a siNA molecule, and (b) assaying the siNA molecule of step (a) under conditions suitable for isolating siNA molecules having improved RNAi activity against the DNA target, such as a gene, chromosome, or portion thereof.

5

10

15

20

25

In one embodiment, the invention features siNA constructs that mediate RNAi in a cell or reconstituted system, wherein the siNA construct comprises one or more chemical modifications described herein that modulates the cellular uptake of the siNA construct.

In another embodiment, the invention features a method for generating siNA molecules against a target gene with improved cellular uptake comprising (a) introducing nucleotides having any of Formula I-VII or any combination thereof into a siNA molecule, and (b) assaying the siNA molecule of step (a) under conditions suitable for isolating siNA molecules having improved cellular uptake.

In one embodiment, the invention features siNA constructs that mediate RNAi against a target gene, wherein the siNA construct comprises one or more chemical modifications described herein that increases the bioavailability of the siNA construct, for example, by attaching polymeric conjugates such as polyethyleneglycol or equivalent conjugates that improve the pharmacokinetics of the siNA construct, or by attaching conjugates that target specific tissue types or cell types *in vivo*. Non-limiting examples of such conjugates are described in Vargeese *et al.*, U.S. Serial No. 10/201,394 incorporated by reference herein.

In one embodiment, the invention features a method for generating siNA molecules of the invention with improved bioavailability, comprising (a) introducing a conjugate into the structure of a siNA molecule, and (b) assaying the siNA molecule of step (a) under conditions suitable for isolating siNA molecules having improved bioavailability. Such conjugates can include ligands for cellular receptors, such as peptides derived from naturally occurring protein ligands; protein localization sequences, including cellular ZIP code sequences; antibodies; nucleic acid aptamers; vitamins and other co-factors, such as folate and N-acetylgalactosamine; polymers, such as polyethyleneglycol (PEG); phospholipids; polyamines, such as spermine or spermidine; and others.

In another embodiment, the invention features a method for generating siNA molecules of the invention with improved bioavailability comprising (a) introducing an excipient formulation to a siNA molecule, and (b) assaying the siNA molecule of step (a) under conditions suitable for isolating siNA molecules having improved bioavailability. Such excipients include polymers such as cyclodextrins, lipids, cationic lipids, polyamines, phospholipids, and others.

5

10

15

20

25

In another embodiment, the invention features a method for generating siNA molecules of the invention with improved bioavailability comprising (a) introducing nucleotides having any of Formulae I-VII or any combination thereof into a siNA molecule, and (b) assaying the siNA molecule of step (a) under conditions suitable for isolating siNA molecules having improved bioavailability.

In another embodiment, polyethylene glycol (PEG) can be covalently attached to siNA compounds of the present invention. The attached PEG can be any molecular weight, preferably from about 2,000 to about 50,000 daltons (Da).

The present invention can be used alone or as a component of a kit having at least one of the reagents necessary to carry out the *in vitro* or *in vivo* introduction of RNA to test samples and/or subjects. For example, preferred components of the kit include a siNA molecule of the invention and a vehicle that promotes introduction of the siNA into cells of interest as described herein (e.g., using lipids and other methods of transfection known in the art, see for example Beigelman *et al*, US 6,395,713). The kit can be used for target validation, such as in determining gene function and/or activity, or in drug optimization, and in drug discovery (see for example Usman et al., USSN 60/402,996). Such a kit can also include instructions to allow a user of the kit to practice the invention.

The term "short interfering nucleic acid", "siNA", "short interfering RNA", "siRNA", "short interfering nucleic acid molecule", "short interfering oligonucleotide molecule", or "chemically-modified short interfering nucleic acid molecule" as used herein refers to any nucleic acid molecule capable of inhibiting or down regulating gene expression or viral replication, for example by mediating RNA interference "RNAi" or gene silencing in a sequence-specific manner; see for example Bass, 2001, *Nature*, 411, 428-429; Elbashir *et al.*, 2001, *Nature*, 411, 494-498; and Kreutzer *et al.*, International PCT Publication No. WO 00/44895; Zernicka-Goetz *et al.*, International PCT Publication No. WO 99/32619; Plaetinck *et al.*, International PCT Publication No. WO 01/29058; Deschamps-Depaillette, International PCT Publication No. WO 99/07409; and Li *et al.*, International PCT Publication No. WO 00/44914; Allshire, 2002, *Science*, 297, 1818-1819; Volpe *et al.*, 2002, *Science*, 297, 1833-1837; Jenuwein, 2002, *Science*, 297, 2215-2218; and Hall *et al.*, 2002, *Science*, 297, 2232-2237;

5

10

15

20

25

PCT/US03/05028 WO 03/074654

Hutvagner and Zamore, 2002, Science, 297, 2056-60; McManus et al., 2002, RNA, 8, 842-850; Reinhart et al., 2002, Gene & Dev., 16, 1616-1626; and Reinhart & Bartel, 2002, Science, 297, 1831). Non limiting examples of siNA molecules of the invention are shown in Figures 4-6, and Tables II, III, and IV herein. For example the siNA can be a double-stranded polynucleotide molecule comprising self-complementary sense and antisense regions, wherein the antisense region comprises nucleotide sequence that is complementary to nucleotide sequence in a target nucleic acid molecule or a portion thereof and the sense region having nucleotide sequence corresponding to the target nucleic acid sequence or a portion thereof. The siNA can be assembled from two separate oligonucleotides, where one strand is the sense strand and the other is the antisense strand, wherein the antisense and sense strands are self-complementary (i.e. each strand comprises nucleotide sequence that is complementary to nucleotide sequence in the other strand; such as where the antisense strand and sense strand form a duplex or double stranded structure, for example wherein the double stranded region is about 19 base pairs); the antisense strand comprises nucleotide sequence that is complementary to nucleotide sequence in a target nucleic acid molecule or a portion thereof and the sense strand comprises nucleotide sequence corresponding to the target nucleic acid sequence or a portion thereof. Alternatively, the siNA is assembled from a single oligonucleotide, where the self-complementary sense and antisense regions of the siNA are linked by means of a nucleic acid based or non-nucleic acid-based linker(s). The siNA can be a polynucleotide with a hairpin secondary structure, having self-complementary sense and antisense regions, wherein the antisense region comprises nucleotide sequence that is complementary to nucleotide sequence in a separate target nucleic acid molecule or a portion thereof and the sense region having nucleotide sequence corresponding to the target nucleic acid sequence or a portion thereof. The siNA can be a circular singlestranded polynucleotide having two or more loop structures and a stem comprising selfcomplementary sense and antisense regions, wherein the antisense region comprises nucleotide sequence that is complementary to nucleotide sequence in a target nucleic acid molecule or a portion thereof and the sense region having nucleotide sequence corresponding to the target nucleic acid sequence or a portion thereof, and wherein the 30 circular polynucleotide can be processed either in vivo or in vitro to generate an active siNA molecule capable of mediating RNAi. The siNA can also comprise a single stranded polynucleotide having nucleotide sequence complementary to nucleotide

5

10

15

20

sequence in a target nucleic acid molecule or a portion thereof (for example, where such siNA molecule does not require the presence within the siNA molecule of nucleotide sequence corresponding to the target nucleic acid sequence or a portion thereof), wherein the single stranded polynucleotide can further comprise a terminal phosphate group, such as a 5'-phosphate (see for example Martinez et al., 2002, Cell., 110, 563-574 and Schwarz et al., 2002, Molecular Cell, 10, 537-568), or 5',3'-diphosphate. In certain embodiment, the siNA molecule of the invention comprises separate sense and antisense sequences or regions, wherein the sense and antisense regions are covalently linked by nucleotide or non-nucleotide linkers molecules as is known in the art, or are alternately non-covalently linked by ionic interactions, hydrogen bonding, van der waals interactions, hydrophobic intercations, and/or stacking interactions. In certain embodiments, the siNA molecules of the invention comprise nucleotide sequence that is complementary to nucleotide sequence of a target gene. In another embodiment, the siNA molecule of the invention interacts with nucleotide sequence of a target gene in a manner that causes inhibition of expression of the target gene. As used herein, siNA molecules need not be limited to those molecules containing only RNA, but further encompasses chemically-modified nucleotides and non-nucleotides. In certain embodiments, the short interfering nucleic acid molecules of the invention lack 2'hydroxy (2'-OH) containing nucleotides. Applicant describes in certain embodiments short interfering nucleic acids that do not require the presence of nucleotides having a 2'hydroxy group for mediating RNAi and as such, short interfering nucleic acid molecules of the invention optionally do not include any ribonucleotides (e.g., nucleotides having a 2'-OH group). Such siNA molecules that do not require the presence of ribonucleotides within the siNA molecule to support RNAi can however have an attached linker or linkers or other attached or associated groups, moieties, or chains containing one or more nucleotides with 2'-OH groups. Optionally, siNA molecules can comprise ribonucleotides at about 5, 10, 20, 30, 40, or 50% of the nucleotide positions. The modified short interfering nucleic acid molecules of the invention can also be referred to as short interfering modified oligonucleotides "siMON." As used herein, the term siNA is meant to be equivalent to other terms used to describe nucleic acid molecules that are capable of mediating sequence specific RNAi, for example short interfering RNA (siRNA), doublestranded RNA (dsRNA), micro-RNA (miRNA), short hairpin RNA (shRNA), short interfering oligonucleotide, short interfering nucleic acid, short interfering modified

5

10

15

20

25

oligonucleotide, chemically-modified siRNA, post-transcriptional gene silencing RNA (ptgsRNA), and others. In addition, as used herein, the term RNAi is meant to be equivalent to other terms used to describe sequence specific RNA interference, such as post transcriptional gene silencing, or epigenetics. For example, siNA molecules of the invention can be used to epigenetically silence genes at both the post-transcriptional level or the pre-transcriptional level. In a non-limiting example, epigenetic regulation of gene expression by siNA molecules of the invention can result from siNA mediated modification of chromatin structure to alter gene expression (see, for example, Allshire, 2002, Science, 297, 1818-1819; Volpe et al., 2002, Science, 297, 1833-1837; Jenuwein, 2002, Science, 297, 2215-2218; and Hall et al., 2002, Science, 297, 2232-2237).

By "modulate" is meant that the expression of the gene, or level of RNA molecule or equivalent RNA molecules encoding one or more proteins or protein subunits, or activity of one or more proteins or protein subunits is up regulated or down regulated, such that expression, level, or activity is greater than or less than that observed in the absence of the modulator. For example, the term "modulate" can mean "inhibit," but the use of the word "modulate" is not limited to this definition.

By "inhibit" it is meant that the activity of a gene expression product or level of RNAs or equivalent RNAs encoding one or more gene products is reduced below that observed in the absence of the nucleic acid molecule of the invention. In one embodiment, inhibition with a siNA molecule preferably is below that level observed in the presence of an inactive or attenuated molecule that is unable to mediate an RNAi response. In another embodiment, inhibition of gene expression with the siNA molecule of the instant invention is greater in the presence of the siNA molecule than in its absence.

By "inhibit", "down-regulate", or "reduce", it is meant that the expression of the gene, or level of RNA molecules or equivalent RNA molecules encoding one or more proteins or protein subunits, or activity of one or more proteins or protein subunits, is reduced below that observed in the absence of the nucleic acid molecules (e.g., siNA) of the invention. In one embodiment, inhibition, down-regulation or reduction with an siNA molecule is below that level observed in the presence of an inactive or attenuated molecule. In another embodiment, inhibition, down-regulation, or reduction with siNA

5

10

15

20

25

molecules is below that level observed in the presence of, for example, an siNA molecule with scrambled sequence or with mismatches. In another embodiment, inhibition, down-regulation, or reduction of gene expression with a nucleic acid molecule of the instant invention is greater in the presence of the nucleic acid molecule than in its absence.

By "gene" or "target gene" is meant, a nucleic acid that encodes an RNA, for example, nucleic acid sequences including, but not limited to, structural genes encoding a polypeptide. The target gene can be a gene derived from a cell, an endogenous gene, a transgene, or exogenous genes such as genes of a pathogen, for example a virus, which is present in the cell after infection thereof. The cell containing the target gene can be derived from or contained in any organism, for example a plant, animal, protozoan, virus, bacterium, or fungus. Non-limiting examples of plants include monocots, dicots, or gymnosperms. Non-limiting examples of animals include vertebrates or invertebrates. Non-limiting examples of fungi include molds or yeasts.

By "endogenous" or "cellular" gene is meant a gene normally found in a cell in its natural location in the genome. For example, HER-2, VEGF, VEGF-R, EGFR, BCL-2, c-MYC, RAS and the like would be considered an endogenous gene. Genes expressed in a cell from a plasmid, viral vector or other vectors or from virus, bacteria, fungi would be considered "foreign" or "heterologous" gene; such genes are not normally found in the host cell, but are introduced by standard gene transfer techniques or as a result of infection by a virus, bacterial or other infectious agent.

By "gene family" is meant a group of more than one nucleic acid molecules that share at least one common characteristic, such as sequence homology, target specificity, mode of action, secondary structure, or the ability to modulate a process or more than one process in a biological system. The gene family can be of viral or cellular origin. The gene family can encode, for example, groups of cytokines, receptors, growth factors, adapter proteins, structural proteins, and other protein epitopes.

By "protein family" is meant a group of more than one proteins, peptides, or polypeptides that share at least one common characteristic, such as sequence homology, target specificity, mode of action, secondary structure, or the ability to modulate a process or more than one process in a biological system. The protein family can be of viral or

5

10

15

20

25

cellular origin. The protein family can encode, for example, groups of cytokines, receptors, growth factors, adapter proteins, structural proteins, and other protein epitopes.

By "highly conserved sequence region" is meant, a nucleotide sequence of one or more regions in a target gene does not vary significantly from one generation to the other or from one biological system to the other.

By "cancer" is meant a group of diseases characterized by uncontrolled growth and spread of abnormal cells.

By "sense region" is meant a nucleotide sequence of a siNA molecule having complementarity to an antisense region of the siNA molecule. In addition, the sense region of a siNA molecule can comprise a nucleic acid sequence having homology with a target nucleic acid sequence.

By "antisense region" is meant a nucleotide sequence of a siNA molecule having complementarity to a target nucleic acid sequence. In addition, the antisense region of a siNA molecule can optionally comprise a nucleic acid sequence having complementarity to a sense region of the siNA molecule.

By "target nucleic acid" is meant any nucleic acid sequence whose expression or activity is to be modulated. The target nucleic acid can be DNA or RNA.

By "complementarity" is meant that a nucleic acid can form hydrogen bond(s) with another nucleic acid sequence by either traditional Watson-Crick or other non-traditional types. In reference to the nucleic molecules of the present invention, the binding free energy for a nucleic acid molecule with its complementary sequence is sufficient to allow the relevant function of the nucleic acid to proceed, e.g., RNAi activity. Determination of binding free energies for nucleic acid molecules is well known in the art (see, e.g., Turner et al., 1987, CSH Symp. Quant. Biol. LII pp.123-133; Frier et al., 1986, Proc. Nat. Acad. Sci. USA 83:9373-9377; Turner et al., 1987, J. Am. Chem. Soc. 109:3783-3785). A percent complementarity indicates the percentage of contiguous residues in a nucleic acid molecule that can form hydrogen bonds (e.g., Watson-Crick base pairing) with a second nucleic acid sequence (e.g., 5, 6, 7, 8, 9, 10 out of 10 being 50%, 60%, 70%, 80%, 90%, and 100% complementary). "Perfectly complementary" means that all the contiguous

5

10

15

20

residues of a nucleic acid sequence will hydrogen bond with the same number of contiguous residues in a second nucleic acid sequence.

The siNA molecules of the invention represent a novel therapeutic approach to a broad spectrum of diseases and conditions, including cancer or cancerous disease, infectious disease, cardiovascular disease, neurological disease, prion disease, inflammatory disease, autoimmune disease, pulmonary disease, renal disease, liver disease, mitochondrial disease, endocrine disease, reproduction related diseases and conditions, and any other indications that can respond to the level of an expressed gene product in a cell or organsim.

In one embodiment of the present invention, each sequence of a siNA molecule of the invention is independently about 18 to about 24 nucleotides in length, in specific embodiments about 18, 19, 20, 21, 22, 23, or 24 nucleotides in length. In another embodiment, the siNA duplexes of the invention independently comprise about 17 to about 23 base pairs (e.g., about 17, 18, 19, 20, 21, 22 or 23). In yet another embodiment, siNA molecules of the invention comprising hairpin or circular structures are about 35 to about 55 (e.g., about 35, 40, 45, 50 or 55) nucleotides in length, or about 38 to about 44 (e.g., 38, 39, 40, 41, 42, 43 or 44) nucleotides in length and comprising about 16 to about 22 (e.g., about 16, 17, 18, 19, 20, 21 or 22) base pairs. Exemplary siNA molecules of the invention are shown in **Table II**. Exemplary synthetic siNA molecules of the invention are shown in **Table II** and/or **Figures 18-19**.

As used herein "cell" is used in its usual biological sense, and does not refer to an entire multicellular organism, e.g., specifically does not refer to a human. The cell can be present in an organism, e.g., birds, plants and mammals such as humans, cows, sheep, apes, monkeys, swine, dogs, and cats. The cell can be prokaryotic or eukaryotic (e.g., mammalian or plant cell). The cell can be of somatic or germ line origin, totipotent or pluripotent, dividing or non-dividing. The cell can also be derived from or can comprise a gamete or embryo, a stem cell, or a fully differentiated cell.

The siNA molecules of the invention are added directly, or can be complexed with cationic lipids, packaged within liposomes, or otherwise delivered to target cells or tissues. The nucleic acid or nucleic acid complexes can be locally administered to relevant tissues ex vivo, or in vivo through injection, infusion pump or stent, with or

5

10

15

20

25

without their incorporation in biopolymers. In particular embodiments, the nucleic acid molecules of the invention comprise sequences shown in **Tables I-II** and/or **Figures 18-19**. Examples of such nucleic acid molecules consist essentially of sequences defined in these tables and figures. Furthermore, the chemically modified constructs described in **Table IV** can be applied to any siNA sequence of the invention.

5

10

15

20

25

BNSDOCID: <WO\_

\_03074654A2\_l\_>

In another aspect, the invention provides mammalian cells containing one or more siNA molecules of this invention. The one or more siNA molecules can independently be targeted to the same or different sites.

By "RNA" is meant a molecule comprising at least one ribonucleotide residue. By "ribonucleotide" is meant a nucleotide with a hydroxyl group at the 2' position of a β-D-ribo-furanose moiety. The terms include double-stranded RNA, single-stranded RNA, isolated RNA such as partially purified RNA, essentially pure RNA, synthetic RNA, recombinantly produced RNA, as well as altered RNA that differs from naturally occurring RNA by the addition, deletion, substitution and/or alteration of one or more nucleotides. Such alterations can include addition of non-nucleotide material, such as to the end(s) of the siNA or internally, for example at one or more nucleotides of the RNA. Nucleotides in the RNA molecules of the instant invention can also comprise non-standard nucleotides, such as non-naturally occurring nucleotides or chemically synthesized nucleotides or deoxynucleotides. These altered RNAs can be referred to as analogs or analogs of naturally-occurring RNA.

By "subject" is meant an organism, which is a donor or recipient of explanted cells or the cells themselves. "Subject" also refers to an organism to which the nucleic acid molecules of the invention can be administered. In one embodiment, a subject is a mammal or mammalian cells. In another embodiment, a subject is a human or human cells.

The term "phosphorothioate" as used herein refers to an internucleotide linkage having Formula I, wherein Z and/or W comprise a sulfur atom. Hence, the term phosphorothioate refers to both phosphorothioate and phosphorodithioate internucleotide linkages.

The term "universal base" as used herein refers to nucleotide base analogs that form base pairs with each of the natural DNA/RNA bases with little discrimination between them. Non-limiting examples of universal bases include C-phenyl, C-naphthyl and other aromatic derivatives, inosine, azole carboxamides, and nitroazole derivatives such as 3-nitropyrrole, 4-nitroindole, 5-nitroindole, and 6-nitroindole as known in the art (see for example Loakes, 2001, *Nucleic Acids Research*, 29, 2437-2447).

The term "acyclic nucleotide" as used herein refers to any nucleotide having an acyclic ribose sugar, for example where any of the ribose carbons (C1, C2, C3, C4, or C5), are independently or in combination absent from the nucleotide.

The nucleic acid molecules of the instant invention, individually, or in combination or in conjunction with other drugs, can be used to treat diseases or conditions discussed herein. For example, to treat a particular disease or condition, the siNA molecules can be administered to a subject or can be administered to other appropriate cells evident to those skilled in the art, individually or in combination with one or more drugs under conditions suitable for the treatment.

In a further embodiment, the siNA molecules can be used in combination with other known treatments to treat conditions or diseases discussed above. For example, the described molecules could be used in combination with one or more known therapeutic agents to treat a disease or condition. Non-limiting examples of other therapeutic agents that can be readily combined with a siNA molecule of the invention are enzymatic nucleic acid molecules, allosteric nucleic acid molecules, antisense, decoy, or aptamer nucleic acid molecules, antibodies such as monoclonal antibodies, small molecules, and other organic and/or inorganic compounds including metals, salts and ions.

In one embodiment, the invention features an expression vector comprising a nucleic acid sequence encoding at least one siNA molecule of the invention, in a manner which allows expression of the siNA molecule. For example, the vector can contain sequence(s) encoding both strands of a siNA molecule comprising a duplex. The vector can also contain sequence(s) encoding a single nucleic acid molecule that is self-complementary and thus forms a siNA molecule. Non-limiting examples of such expression vectors are described in Paul et al., 2002, Nature Biotechnology, 19, 505; Miyagishi and Taira, 2002, Nature Biotechnology, 19, 497; Lee et al., 2002, Nature

5

10

15

20

25

Biotechnology, 19, 500; and Novina et al., 2002, Nature Medicine, advance online publication doi:10.1038/nm725.

In another embodiment, the invention features a mammalian cell, for example, a human cell, including an expression vector of the invention.

In yet another embodiment, the expression vector of the invention comprises a sequence for a siRNA molecule having complementarity to a RNA molecule referred to by a Genbank Accession number in Table III.

In yet another embodiment, the expression vector of the invention comprises a sequence for a siNA molecule having complementarity to a RNA molecule referred to by a Genbank Accession numbers, for example Genbank Accession Nos. shown in **Table I**.

In one embodiment, an expression vector of the invention comprises a nucleic acid sequence encoding two or more siNA molecules, which can be the same or different.

In another aspect of the invention, siRNA molecules that interact with target RNA molecules and down-regulate gene encoding target RNA molecules (for example target RNA molecules referred to by Genbank Accession number in Table III) are expressed The recombinant vectors from transcription units inserted into DNA or RNA vectors. can be DNA plasmids or viral vectors. siNA expressing viral vectors can be constructed based on, but not limited to, adeno-associated virus, retrovirus, adenovirus, or alphavirus. The recombinant vectors capable of expressing the siNA molecules can be delivered as described herein, and persist in target cells. Alternatively, viral vectors can be used that provide for transient expression of siNA molecules. Such vectors can be repeatedly administered as necessary. Once expressed, the siNA molecules bind and down-regulate gene function or expression via RNA interference (RNAi). Delivery of siNA expressing vectors can be systemic, such as by intravenous or intramuscular administration, by administration to target cells ex-planted from a subject followed by reintroduction into the subject, or by any other means that would allow for introduction into the desired target cell.

By "vectors" is meant any nucleic acid- and/or viral-based technique used to deliver a desired nucleic acid.

5

10

15

20

Other features and advantages of the invention will be apparent from the following description of the preferred embodiments thereof, and from the claims.

## BRIEF DESCRIPTION OF THE DRAWINGS

Figure 1 shows a non-limiting example of a scheme for the synthesis of siNA molecules. The complementary siNA sequence strands, strand 1 and strand 2, are synthesized in tandem and are connected by a cleavable linkage, such as a nucleotide succinate or abasic succinate, which can be the same or different from the cleavable linker used for solid phase synthesis on a solid support. The synthesis can be either solid phase or solution phase, in the example shown, the synthesis is a solid phase synthesis. The synthesis is performed such that a protecting group, such as a dimethoxytrityl group, remains intact on the terminal nucleotide of the tandem oligonucleotide. Upon cleavage and deprotection of the oligonucleotide, the two siNA strands spontaneously hybridize to form a siNA duplex, which allows the purification of the duplex by utilizing the properties of the terminal protecting group, for example by applying a trityl on purification method wherein only duplexes/oligonucleotides with the terminal protecting group are isolated.

Figure 2 shows a MALDI-TOV mass spectrum of a purified siNA duplex synthesized by a method of the invention. The two peaks shown correspond to the predicted mass of the separate siNA sequence strands. This result demonstrates that the siNA duplex generated from tandem synthesis can be purified as a single entity using a simple trityl-on purification methodology.

Figure 3 shows the results of a stability assay used to determine the serum stability of chemically modified siNA constructs compared to a siNA control consisting of all RNA with 3'-TT termini. T ½ values are shown for duplex stability.

Figure 4 shows the results of an RNAi activity screen of phosphorothioate modified siNA constructs using a luciferase reporter system.

Figure 5 shows the results of an RNAi activity screen of phosphorothioate and universal base modified siNA constructs using a luciferase reporter system.

5

10

15

20

Figure 6 shows the results of an RNAi activity screen of 2'-O-methyl modified siNA constructs using a luciferase reporter system.

Figure 7 shows the results of an RNAi activity screen of 2'-O-methyl and 2'-deoxy-2'-fluoro modified siNA constructs using a luciferase reporter system.

Figure 8 shows the results of an RNAi activity screen of a phosphorothicate modified siNA construct using a luciferase reporter system.

Figure 9 shows the results of an RNAi activity screen of an inverted deoxyabasic modified siNA construct generated via tandem synthesis using a luciferase reporter system.

Figure 10 shows the results of an RNAi activity screen of chemically modified siNA constructs including 3'-glyceryl modified siNA constructs compared to an all RNA control siNA construct using a luciferase reporter system. These chemically modified siNAs were compared in the luciferase assay described herein at 1 nM and 10nM concentration using an all RNA siNA control (siGL2) having having 3'-terminal dithymidine (TT) and its corresponding inverted control (Inv siGL2). The background level of luciferase expression in the HeLa cells is designated by the "cells" column. Sense and antisense strands of chemically modified siNA constructs are shown by RPI number (sense strand/antisense strand). Sequences correspoding to these RPI numbers are shown in Table I.

Figure 11 shows the results of an RNAi activity screen of chemically modified siNA constructs. The screen compared various combinations of sense strand chemical modifications and antisense strand chemical modifications. These chemically modified siNAs were compared in the luciferase assay described herein at 1 nM and 10nM concentration using an all RNA siNA control (siGL2) having having 3'-terminal dithymidine (TT) and its corresponding inverted control (Inv siGL2). The background level of luciferase expression in the HeLa cells is designated by the "cells" column. Sense and antisense strands of chemically modified siNA constructs are shown by RPI number (sense strand/antisense strand). Sequences correspoding to these RPI numbers are shown in Table I.

5

10

15

20

Figure 12 shows the results of an RNAi activity screen of chemically modified siNA constructs. The screen compared various combinations of sense strand chemical modifications and antisense strand chemical modifications. These chemically modified siNAs were compared in the luciferase assay described herein at 1 nM and 10nM concentration using an all RNA siNA control (siGL2) having having 3'-terminal dithymidine (TT) and its corresponding inverted control (Inv siGL2). The background level of luciferase expression in the HeLa cells is designated by the "cells" column. Sense and antisense strands of chemically modified siNA constructs are shown by RPI number (sense strand/antisense strand). Sequences correspoding to these RPI numbers are shown in Table I. In addition, the antisense strand alone (RPI 30430) and an inverted control (RPI 30227/30229, having matched chemistry to RPI 30063/30224) was compared to the siNA duplexes described above.

Figure 13 shows the results of an RNAi activity screen of chemically modified siNA constructs. The screen compared various combinations of sense strand chemical modifications and antisense strand chemical modifications. These chemically modified siNAs were compared in the luciferase assay described herein at 1 nM and 10nM concentration using an all RNA siNA control (siGL2) having having 3'-terminal dithymidine (TT) and its corresponding inverted control (Inv siGL2). The background level of luciferase expression in the HeLa cells is designated by the "cells" column. Sense and antisense strands of chemically modified siNA constructs are shown by RPI number (sense strand/antisense strand). Sequences correspoding to these RPI numbers are shown in Table I. In addition, an inverted control (RPI 30226/30229, having matched chemistry to RPI 30222/30224) was compared to the siNA duplexes described above.

Figure 14 shows the results of an RNAi activity screen of chemically modified siNA constructs including various 3'-terminal modified siNA constructs compared to an all RNA control siNA construct using a luciferase reporter system. These chemically modified siNAs were compared in the luciferase assay described herein at 1 nM and 10nM concentration using an all RNA siNA control (siGL2) having having 3'-terminal dithymidine (TT) and its corresponding inverted control (Inv siGL2). The background level of luciferase expression in the HeLa cells is designated by the "cells" column. Sense and antisense strands of chemically modified siNA constructs are shown by RPI

5

10

15

20

25

number (sense strand/antisense strand). Sequences correspoding to these RPI numbers are shown in Table I.

Figure 15 shows the results of an RNAi activity screen of chemically modified siNA constructs. The screen compared various combinations of sense strand chemistries compared to a fixed antisense strand chemistry. These chemically modified siNAs were compared in the luciferase assay described herein at 1 nM and 10nM concentration using an all RNA siNA control (siGL2) having having 3'-terminal dithymidine (TT) and its corresponding inverted control (Inv siGL2). The background level of luciferase expression in the HeLa cells is designated by the "cells" column. Sense and antisense strands of chemically modified siNA constructs are shown by RPI number (sense strand/antisense strand). Sequences corresponding to these RPI numbers are shown in Table I.

Figure 16 shows the results of a siNA titration study wherein the RNAi activity of a phosphorothicate modified siNA construct is compared to that of a siNA construct consisting of all ribonucleotides except for two terminal thymidine residues using a luciferase reporter system.

Figure 17 shows a non-limiting proposed mechanistic representation of target RNA degradation involved in RNAi. Double-stranded RNA (dsRNA), which is generated by RNA-dependent RNA polymerase (RdRP) from foreign single-stranded RNA, for example viral, transposon, or other exogenous RNA, activates the DICER enzyme that in turn generates siNA duplexes. Alternately, synthetic or expressed siNA can be introduced directely into a cell by appropriate means. An active siNA complex forms which recognizes a target RNA, resulting in degradation of the target RNA by the RISC endonuclease complex or in the synthesis of additional RNA by RNA-dependent RNA polymerase (RdRP), which can activate DICER and result in additional siNA molecules, thereby amplifying the RNAi response.

Figure 18A-F shows non-limiting examples of chemically-modified siNA constructs of the present invention. In the figure, N stands for any nucleotide (adenosine, guanosine, cytosine, uridine, or optionally thymidine, for example thymidine can be substituted in the overhanging regions designated by parenthesis (N N). Various modifications are shown for the sense and antisense strands of the siNA constructs.

5

10

15

20

25

The sense strand comprises 21 nucleotides having four Figure 18A: phosphorothioate 5'- and 3'-terminal internucleotide linkages, wherein the two terminal 3'nucleotides are optionally base paired and wherein all pyrimidine nucleotides that may be present are 2'-O-methyl or 2'-deoxy-2'-fluoro modified nucleotides except for (N N) nucleotides, which can comprise ribonucleotides, deoxynucleotides, universal bases, or other chemical modifications described herein. The antisense strand comprises 21 nucleotides, optionally having a 3'-terminal glyceryl moiety and wherein the two terminal 3'-nucleotides are optionally complementary to the target RNA sequence, and having one phosphorothioate internucleotide linkage and four 5'-terminal 3'-terminal phosphorothioate internucleotide linkages and wherein all pyrimidine nucleotides that may be present are 2'-deoxy-2'-fluoro modified nucleotides except for (N N) nucleotides, which can comprise ribonucleotides, deoxynucleotides, universal bases, or other chemical modifications described herein.

Figure 18B: The sense strand comprises 21 nucleotides wherein the two terminal 3'-nucleotides are optionally base paired and wherein all pyrimidine nucleotides that may be present are 2'-O-methyl or 2'-deoxy-2'-fluoro modified nucleotides except for (N N) nucleotides, which can comprise ribonucleotides, deoxynucleotides, universal bases, or other chemical modifications described herein. The antisense strand comprises 21 nucleotides, optionally having a 3'-terminal glyceryl moiety and wherein the two terminal 3'-nucleotides are optionally complementary to the target RNA sequence, and wherein all pyrimidine nucleotides that may be present are 2'-deoxy-2'-fluoro modified nucleotides except for (N N) nucleotides, which can comprise ribonucleotides, deoxynucleotides, universal bases, or other chemical modifications described herein.

Figure 18C: The sense strand comprises 21 nucleotides having 5'- and 3'- terminal cap moieties wherein the two terminal 3'-nucleotides are optionally base paired and wherein all pyrimidine nucleotides that may be present are 2'-O-methyl or 2'-deoxy-2'-fluoro modified nucleotides except for (N N) nucleotides, which can comprise ribonucleotides, deoxynucleotides, universal bases, or other chemical modifications described herein. The antisense strand comprises 21 nucleotides, optionally having a 3'-terminal glyceryl moiety and wherein the two terminal 3'-nucleotides are optionally complementary to the target RNA sequence, and having one 3'-terminal phosphorothioate internucleotide linkage and wherein all pyrimidine nucleotides that may be present are 2'-

5

10

15

20

25

deoxy-2'-fluoro modified nucleotides except for (N N) nucleotides, which can comprise ribonucleotides, deoxynucleotides, universal bases, or other chemical modifications described herein.

Figure 18D: The sense strand comprises 21 nucleotides having 5'- and 3'- terminal cap moieties wherein the two terminal 3'-nucleotides are optionally base paired and wherein all pyrimidine nucleotides that may be present are 2'-deoxy-2'-fluoro modified nucleotides except for (N N) nucleotides, which can comprise ribonucleotides, deoxynucleotides, universal bases, or other chemical modifications described herein and wherein and all purine nucleotides that may be present are 2'-deoxy nucleotides. The antisense strand comprises 21 nucleotides, optionally having a 3'-terminal glyceryl moiety and wherein the two terminal 3'-nucleotides are optionally complementary to the target RNA sequence, and having one 3'-terminal phosphorothioate internucleotide linkage and wherein all pyrimidine nucleotides that may be present are 2'-deoxy-2'-fluoro modified nucleotides and all purine nucleotides that may be present are 2'-O-methyl modified nucleotides except for (N N) nucleotides, which can comprise ribonucleotides, deoxynucleotides, universal bases, or other chemical modifications described herein.

Figure 18E: The sense strand comprises 21 nucleotides having 5'- and 3'- terminal cap moieties wherein the two terminal 3'-nucleotides are optionally base paired and wherein all pyrimidine nucleotides that may be present are 2'-deoxy-2'-fluoro modified nucleotides except for (N N) nucleotides, which can comprise ribonucleotides, deoxynucleotides, universal bases, or other chemical modifications described herein. The antisense strand comprises 21 nucleotides, optionally having a 3'-terminal glyceryl moiety and wherein the two terminal 3'-nucleotides are optionally complementary to the target RNA sequence, and wherein all pyrimidine nucleotides that may be present are 2'-deoxy-2'-fluoro modified nucleotides and all purine nucleotides that may be present are 2'-O-methyl modified nucleotides except for (N N) nucleotides, which can comprise ribonucleotides, deoxynucleotides, universal bases, or other chemical modifications described herein.

Figure 18F: The sense strand comprises 21 nucleotides having 5'- and 3'- terminal cap moieties wherein the two terminal 3'-nucleotides are optionally base paired and wherein all pyrimidine nucleotides that may be present are 2'-deoxy-2'-fluoro modified

5

10

15

20

25

nucleotides except for (N N) nucleotides, which can comprise ribonucleotides, deoxynucleotides, universal bases, or other chemical modifications described herein. The antisense strand comprises 21 nucleotides, optionally having a 3'-terminal glyceryl moiety and wherein the two terminal 3'-nucleotides are optionally complementary to the target RNA sequence, and having one 3'-terminal phosphorothioate internucleotide linkage and wherein all pyrimidine nucleotides that may be present are 2'-deoxy-2'-fluoro modified nucleotides and all purine nucleotides that may be present are 2'-deoxy modified nucleotides except for (N N) nucleotides, which can comprise ribonucleotides, deoxynucleotides, universal bases, or other chemical modifications described herein. The antisense strand of constructs A-F comprise sequence complementary to target RNA sequence of the invention.

Figure 19 shows non-limiting examples of specific chemically modified siNA sequences of the invention. A-F applies the chemical modifications described in Figure 18A-F to a representative siNA sequence targeting the EGFR (HER1).

Figure 20 shows non-limiting examples of different siNA constructs of the invention. The examples shown (constructs 1, 2, and 3) have 19 representative base pairs, however, different embodiments of the invention include any number of base pairs Bracketed regions represent nucleotide overhangs, for example described herein. comprising between about 1, 2, 3, or 4 nucleotides in length, preferably about 2 nucleotides. Constructs 1 and 2 can be used independently for RNAi activity. Construct 2 can comprise a polynucleotide or non-nucleotide linker, which can optionally be designed as a biodegradable linker. In one embodiment, the loop structure shown in construct 2 can comprise a biodegradable linker that results in the formation of construct 1 in vivo and/or in vitro. In another example, construct 3 can be used to generate construct 2 under the same principle wherein a linker is used to generate the active siNA construct 2 in vivo and/or in vitro, which can optionally utilize another biodegradable linker to generate the active siNA construct 1 in vivo and/or in vitro. As such, the stability and/or activity of the siNA constructs can be modulated based on the design of the siNA construct for use in vivo or in vitro and/or in vitro.

Figure 21 is a diagrammatic representation of a method used to determine target sites for siNA mediated RNAi within a particular target nucleic acid sequence, such as

5

10

15

20

25

messenger RNA. (A) A pool of siNA oligonucleotides are synthesized wherein the antisense region of the siNA constructs has complementarity to target sites across the target nucleic acid sequence, and wherein the sense region comprises sequence complementary to the antisense region of the siNA. (B) The sequences are transfected into cells. (C) Cells are selected based on phenotypic change that is associated with modulation of the target nucleic acid sequence. (D) The siNA is isolated from the selected cells and is sequenced to identify efficacious target sites within the target nucleic acid sequence.

Figure 22 shows non-limiting examples of different stabilization chemistries (1-10) that can be used, for example, to stabilize the 3'-end of siNA sequences of the invention, including (1) [3-3']-inverted deoxyribose; (2) deoxyribonucleotide; (3) [5'-3']-3'-deoxyribonucleotide; (4) [5'-3']-ribonucleotide; (5) [5'-3']-3'-O-methyl ribonucleotide; (6) 3'-glyceryl; (7) [3'-5']-3'-deoxyribonucleotide; (8) [3'-3']-deoxyribonucleotide; (9) [5'-2']-deoxyribonucleotide; and (10) [5-3']-dideoxyribonucleotide. In addition to modified and unmodified backbone chemistries indicated in the figure, these chemistries can be combined with different backbone modifications as described herein, for example, backbone modifications having Formula I. In addition, the 2'-deoxy nucleotide shown 5' to the terminal modifications shown can be another modified or unmodified nucleotide or non-nucleotide described herein, for example modifications having any of Formulae I-VII or any combination thereof.

Figure 23 shows a non-limiting example of siNA mediated inhibition of VEGF-induced angiogenesis using the rat corneal model of angiogenesis. siNA targeting site 2340 of VEGFR1 RNA (shown as RPI No. sense strand/antisense strand) were compared to inverted controls (shown as RPI No. sense strand/antisense strand) at three different concentrations and compared to a VEGF control in which no siNA was administered.

Figure 24 shows a non-limiting example of a strategy used to identify chemically modified siNA constructs of the invention that are nuclease resistance while preserving the ability to mediate RNAi activity. Chemical modifications are introduced into the siNA construct based on educated design parameters (e.g. introducing 2'-mofications, base modifications, backbone modifications, terminal cap modifications etc). The modified construct in tested in an appropriate system (e.g human serum for nuclease

5

10

15

20

25

resistance, shown, or an animal model for PK/delivery parameters). In parallel, the siNA construct is tested for RNAi activity, for example in a cell culture system such as a luciferase reporter assay). Lead siNA constructs are then identified which possess a particular characteristic while maintaining RNAi activity, and can be further modified and assayed once again. This same approach can be used to identify siNA-conjugate molecules with improved pharmacokinetic profiles, delivery, and RNAi activity.

Figure 25 shows a non-limiting example of reduction of HER2 mRNA in A549 cells mediated by RNA-based and chemically-modified siNAs that target HER2 mRNA sites 2344 and 3706. A549 cells were transfected with 4 ug/ml lipid complexed with 25 nM unmodified siNA with a 3'-terminal dithymidine cap (RPI#28266/28267) or a corresponding inverted control (RPI#28268/28269) for site 2344 and (RPI#28262/28263) and a corresponding inverted control (RPI 28264/28265) for site 3706. In addition, A549 cells were transfected with 4 ug/ml lipid complexed with 25 nM modified siNA (RPI#30442/30443) and a corresponding matched control (RPI#30444/30445) for site 2344 and (RPI#30438/30439) and a corresponding matched control (RPI 30440/30441) for site 3706. As shown in the figures, the modified and unmodified constructs targeting sites 2344 and 3706 all demonstrate significant inhibition of HER2 RNA expression.

Figure 26 shows a non-limiting example of reduction of PKC-alpha mRNA in A549 cells mediated by chemically-modified siNAs that target PKC-alpha mRNA. A549 cells were transfected with 0.25 ug/well of lipid complexed with 25 nM siNA. A screen of siNA constructs comprising ribonucleotides and 3'-terminal dithymidine caps was compared to untreated cells, scrambled siNA control constructs (Scram1 and Scram2), and cells transfected with lipid alone (transfection control). As shown in the figure, all of the siNA constructs show significant reduction of PKC-alpha RNA expression.

Figure 27 shows a non-limiting example of reduction of Myc (c-Myc) mRNA in 293T cells mediated by chemically-modified siNAs that target c-Myc mRNA. 293T cells were transfected with 0.25 ug/well of lipid complexed with 25 nM siNA. A screen of siNA constructs comprising ribonucleotides and 3'-terminal dithymidine caps was compared to untreated cells, scrambled siNA control constructs (Scram1 and Scram2), and cells transfected with lipid alone (transfection control). As shown in the figure, three

5

10

15

20

25

of the siNA constructs (RPI 30993/31069; RPI 30995/31071; and RPI 30996/31072) show significant reduction of c-Myc RNA expression.

Figure 28 shows a non-limiting example of reduction of BCL2 mRNA in A549 cells mediated by chemically-modified siNAs that target BCL2 mRNA. A549 cells were transfected with 0.25 ug/well of lipid complexed with 25 nM siNA. A siNA construct comprising ribonucleotides and 3'-terminal dithymidine caps (RPI#30998/31074) was tested along with a chemically modified siNA construct comprising 2'-deoxy-2'-fluoro pyrimidine nucleotides and purine ribonucleotides in which the sense strand of the siNA is further modified with 5' and 3'-terminal inverted deoxyabasic caps and the antisense linkage internucleotide phosphorothioate 3'-terminal comprises a strand (RPI#31368/31369), which was also compared to a matched chemistry inverted control (RPI#31370/31371) and a chemically modified siNA construct comprising 2'-deoxy-2'fluoro pyrimidine and 2'-deoxy-2'-fluoro purine nucleotides in which the sense strand of the siNA is further modified with 5' and 3'-terminal inverted deoxyabasic caps and the antisense strand comprises a 3'-terminal phosphorothioate internucleotide linkage (RPI#31372/31373) which was also compared to a matched chemistry inverted control (RPI#31374/31375). In addition, the siNA constructs were also compared to untreated cells, cells transfected with lipid and scrambled siNA constructs (Scram1 and Scram2). and cells transfected with lipid alone (transfection control). As shown in the figure, the siNA constructs show significant reduction of BCL2 RNA expression compared to scrambled, untreated, and transfection controls.

Figure 29 shows a non-limiting example of reduction of CHK-1 mRNA in A549 cells mediated by chemically-modified siNAs that target CHK-1 mRNA. A549 cells were transfected with 0.25 ug/well of lipid complexed with 25 nM siNA. A siNA dithymidine 3'-terminal ribonucleotides and comprising construct (RPI#31003/31079) and a chemically modified siNA construct comprising 2'-deoxy-2'fluoro pyrimidine nucleotides and purine ribonucleotides in which the sense strand of the siNA is further modified with 5' and 3'-terminal inverted deoxyabasic caps and in which the antisense strand comprises a 3'-terminal phosphorothioate internucleotide linkage (RPI#31302/31303), were compared to a matched chemistry inverted control (RPI#31314/31325). In addition, the siNA constructs were also compared to untreated cells, cells transfected with lipid and scrambled siNA constructs (Scram1 and Scram2),

5

10

15

20

25

and cells transfected with lipid alone (transfection control). As shown in the figure, both siNA constructs show significant reduction of CHK-1 RNA expression compared to appropriate controls.

Figure 30 shows a non-limiting example of reduction of BACE mRNA in A549 cells mediated by siNAs that target BACE mRNA. A549 cells were transfected with 0.25 ug/well of lipid complexed with 25 nM siNA. A screen of siNA constructs comprising ribonucleotides and 3'-terminal dithymidine caps was compared to untreated cells, scrambled siNA control constructs (Scram1 and Scram2), and cells transfected with lipid alone (transfection control). As shown in the figure, all of the siNA constructs show significant reduction of BACE RNA expression.

Figure 31 shows a non-limiting example of reduction of cyclin D1 mRNA in A549 cells mediated by chemically-modified siNAs that target cyclin D1 mRNA. A549 cells were transfected with 0.25 ug/well of lipid complexed with 25 nM siNA. A siNA comprising construct ribonucleotides and 3'-terminal dithymidine (RPI#31009/31085) was compared to a chemically modified siNA construct comprising 2'-deoxy-2'-fluoro pyrimidine nucleotides and purine ribonucleotides in which the sense strand of the siNA is further modified with 5' and 3'-terminal inverted deoxyabasic caps and the antisense strand comprises a 3'-terminal phosphorothioate internucleotide linkage (RPI#31304/31305), which was also compared to a matched chemistry inverted control (RPI#31316/31317). In addition, the siNA constructs were also compared to untreated cells, cells transfected with lipid and scrambled siNA constructs (Scram1 and Scram2), and cells transfected with lipid alone (transfection control). As shown in the figure, both siNA constructs show significant reduction of cyclin D1 RNA expression.

Figure 32 shows a non-limiting example of reduction of PTP-1B mRNA in A549 cells mediated by chemically-modified siNAs that target PTP-1B mRNA. A549 cells were transfected with 0.25 ug/well of lipid complexed with 25 nM siNA. A siNA construct comprising ribonucleotides and 3'-terminal dithymidine caps (RPI#31018/31307) was compared to a chemically modified siNA construct comprising 2'-deoxy-2'-fluoro pyrimidine nucleotides and purine ribonucleotides in which the sense strand of the siNA is further modified with 5' and 3'-terminal inverted deoxyabasic caps and the antisense strand comprises a 3'-terminal phosphorothioate internucleotide linkage

5

10

15

20

25

(RPI#31306/31307), which was also compared to a matched chemistry inverted control (RPI#31318/31319). In addition, the siNA constructs were also compared to untreated cells, cells transfected with lipid and scrambled siNA constructs (Scram1 and Scram2), and cells transfected with lipid alone (transfection control). As shown in the figure, both siNA constructs show significant reduction of PTP-1B RNA expression.

Figure 33 shows a non-limiting example of reduction of ERG2 mRNA in DLD1 cells mediated by siNAs that target ERG2 mRNA. DLD1 cells were transfected with 0.25 ug/well of lipid complexed with 25 nM siNA. A screen of siNA constructs comprising ribonucleotides and 3'-terminal dithymidine caps was compared to untreated cells, scrambled siNA control constructs (Scram1 and Scram2), and cells transfected with lipid alone (transfection control). As shown in the figure, all of the siNA constructs show significant reduction of ERG2 RNA expression.

Figure 34 shows a non-limiting example of reduction of PCNA mRNA in A549 cells mediated by chemically-modified siNAs that target PCNA mRNA. A549 cells were transfected with 0.25 ug/well of lipid complexed with 25 nM siNA. A siNA construct comprising ribonucleotides and 3'-terminal dithymidine caps (RPI#31035/31111) was compared to a chemically modified siNA construct comprising 2'-deoxy-2'-fluoro pyrimidine nucleotides and purine ribonucleotides in which the sense strand of the siNA is further modified with 5' and 3'-terminal inverted deoxyabasic caps and the antisense linkage phosphorothioate internucleotide 3'-terminal comprises a strand (RPI#31310/31311), which was also compared to a matched chemistry inverted control (RPI#31322/31323). In addition, the siNA constructs were also compared to untreated cells, cells transfected with lipid and scrambled siNA constructs (Scram1 and Scram2), and cells transfected with lipid alone (transfection control). As shown in the figure, both siNA constructs show significant reduction of PCNA RNA expression.

# DETAILED DESCRIPTION OF THE INVENTION

# Mechanism of action of Nucleic Acid Molecules of the Invention

The discussion that follows discusses the proposed mechanism of RNA interference mediated by short interfering RNA as is presently known, and is not meant to be limiting and is not an admission of prior art. Applicant demonstrates herein that chemically-

5

10

15

20

25

modified short interfering nucleic acids possess similar or improved capacity to mediate RNAi as do siRNA molecules and are expected to possess improved stability and activity in vivo; therefore, this discussion is not meant to be limiting only to siRNA and can be applied to siNA as a whole. By "improved capacity to mediate RNAi" or "improved RNAi activity" is meant to include RNAi activity measured in vitro and/or in vivo where the RNAi activity is a reflection of both the ability of the siNA to mediate RNAi and the stability of the siNAs of the invention. In this invention, the product of these activities can be increased in vitro and/or in vivo compared to an all RNA siRNA or a siNA containing a plurality of ribonucleotides. In some cases, the activity or stability of the siNA molecule can be decreased (i.e., less than ten-fold), but the overall activity of the siNA molecule is enhanced in vitro and/or in vivo.

RNA interference refers to the process of sequence specific post-transcriptional gene silencing in animals mediated by short interfering RNAs (siRNAs) (Fire et al., 1998, Nature, 391, 806). The corresponding process in plants is commonly referred to as posttranscriptional gene silencing or RNA silencing and is also referred to as quelling in The process of post-transcriptional gene silencing is thought to be an evolutionarily-conserved cellular defense mechanism used to prevent the expression of foreign genes which is commonly shared by diverse flora and phyla (Fire et al., 1999, Trends Genet., 15, 358). Such protection from foreign gene expression may have evolved in response to the production of double-stranded RNAs (dsRNAs) derived from viral infection or the random integration of transposon elements into a host genome via a cellular response that specifically destroys homologous single-stranded RNA or viral genomic RNA. The presence of dsRNA in cells triggers the RNAi response though a mechanism that has yet to be fully characterized. This mechanism appears to be different from the interferon response that results from dsRNA-mediated activation of protein kinase PKR and 2', 5'-oligoadenylate synthetase resulting in non-specific cleavage of mRNA by ribonuclease L.

1 1

The presence of long dsRNAs in cells stimulates the activity of a ribonuclease III enzyme referred to as Dicer. Dicer is involved in the processing of the dsRNA into short pieces of dsRNA known as short interfering RNAs (siRNAs) (Berstein *et al.*, 2001, *Nature*, 409, 363). Short interfering RNAs derived from Dicer activity are typically about 21 to about 23 nucleotides in length and comprise about 19 base pair duplexes. Dicer has

5

10

15

20

25

also been implicated in the excision of 21- and 22-nucleotide small temporal RNAs (stRNAs) from precursor RNA of conserved structure that are implicated in translational control (Hutvagner et al., 2001, Science, 293, 834). The RNAi response also features an endonuclease complex containing a siRNA, commonly referred to as an RNA-induced silencing complex (RISC), which mediates cleavage of single-stranded RNA having sequence homologous to the siRNA. Cleavage of the target RNA takes place in the middle of the region complementary to the guide sequence of the siRNA duplex (Elbashir et al., 2001, Genes Dev., 15, 188). In addition, RNA interference can also involve small RNA (e.g., micro-RNA or miRNA) mediated gene silencing, presumably though cellular mechanisms that regulate chromatin structure and thereby prevent transcription of target gene sequences (see for example Allshire, 2002, Science, 297, 1818-1819; Volpe et al., 2002, Science, 297, 1833-1837; Jenuwein, 2002, Science, 297, 2215-2218; and Hall et al., 2002, Science, 297, 2232-2237). As such, siNA molecules of the invention can be used to mediate gene silencing via interaction with RNA transcripts or alternately by interaction with particular gene sequences, wherein such interaction results in gene silencing either at the transcriptional level or post-transcriptional level.

RNAi has been studied in a variety of systems. Fire et al., 1998, Nature, 391, 806, were the first to observe RNAi in C. elegans. Wianny and Goetz, 1999, Nature Cell Biol., 2, 70, describe RNAi mediated by dsRNA in mouse embryos. Hammond et al., 2000, Nature, 404, 293, describe RNAi in Drosophila cells transfected with dsRNA. Elbashir et al., 2001, Nature, 411, 494, describe RNAi induced by introduction of duplexes of synthetic 21-nucleotide RNAs in cultured mammalian cells including human embryonic kidney and HeLa cells. Recent work in Drosophila embryonic lysates has revealed certain requirements for siRNA length, structure, chemical composition, and sequence that are essential to mediate efficient RNAi activity. These studies have shown that 21 nucleotide siRNA duplexes are most active when containing two 2-nucleotide 3'terminal nucleotide overhangs. Furthermore, substitution of one or both siRNA strands with 2'-deoxy or 2'-O-methyl nucleotides abolishes RNAi activity, whereas substitution of 3'-terminal siRNA nucleotides with deoxy nucleotides was shown to be tolerated. Mismatch sequences in the center of the siRNA duplex were also shown to abolish RNAi activity. In addition, these studies also indicate that the position of the cleavage site in the target RNA is defined by the 5'-end of the siRNA guide sequence rather than the 3'-end

5

10

15

20

25

(Elbashir et al., 2001, EMBO J., 20, 6877). Other studies have indicated that a 5'-phosphate on the target-complementary strand of a siRNA duplex is required for siRNA activity and that ATP is utilized to maintain the 5'-phosphate moiety on the siRNA (Nykanen et al., 2001, Cell, 107, 309); however, siRNA molecules lacking a 5'-phosphate are active when introduced exogenously, suggesting that 5'-phosphorylation of siRNA constructs may occur in vivo.

#### Synthesis of Nucleic acid Molecules

5

10

15

20

25

30

Synthesis of nucleic acids greater than 100 nucleotides in length is difficult using automated methods, and the therapeutic cost of such molecules is prohibitive. In this invention, small nucleic acid motifs "small" refers to nucleic acid motifs no more than 100 nucleotides in length, preferably no more than 80 nucleotides in length, and most preferably no more than 50 nucleotides in length; *e.g.*, individual siNA oligonucleotide sequences or siNA sequences synthesized in tandem) are preferably used for exogenous delivery. The simple structure of these molecules increases the ability of the nucleic acid to invade targeted regions of protein and/or RNA structure. Exemplary molecules of the instant invention are chemically synthesized, and others can similarly be synthesized.

Oligonucleotides (e.g., certain modified oligonucleotides or portions of oligonucleotides lacking ribonucleotides) are synthesized using protocols known in the art, for example as described in Caruthers et al., 1992, Methods in Enzymology 211, 3-19, Thompson et al., International PCT Publication No. WO 99/54459, Wincott et al., 1995, Nucleic Acids Res. 23, 2677-2684, Wincott et al., 1997, Methods Mol. Bio., 74, 59, Brennan et al., 1998, Biotechnol Bioeng., 61, 33-45, and Brennan, U.S. Pat. No. 6,001,311. All of these references are incorporated herein by reference. The synthesis of oligonucleotides makes use of common nucleic acid protecting and coupling groups, such as dimethoxytrityl at the 5'-end, and phosphoramidites at the 3'-end. In a non-limiting example, small scale syntheses are conducted on a 394 Applied Biosystems, Inc. synthesizer using a 0.2 µmol scale protocol with a 2.5 min coupling step for 2'-O-methylated nucleotides and a 45 sec coupling step for 2'-deoxy nucleotides or 2'-deoxy-2'-fluoro nucleotides. Table II outlines the amounts and the contact times of the reagents used in the synthesis cycle. Alternatively, syntheses at the 0.2 µmol scale can be performed on a 96-well plate synthesizer, such as the instrument produced by Protogene

(Palo Alto, CA) with minimal modification to the cycle. A 33-fold excess (60  $\mu L$  of 0.11  $M = 6.6 \mu mol$ ) of 2'-O-methyl phosphoramidite and a 105-fold excess of S-ethyl tetrazole (60  $\mu$ L of 0.25 M = 15  $\mu$ mol) can be used in each coupling cycle of 2'-O-methyl residues relative to polymer-bound 5'-hydroxyl. A 22-fold excess (40  $\mu$ L of 0.11 M = 4.4  $\mu$ mol) of deoxy phosphoramidite and a 70-fold excess of S-ethyl tetrazole (40  $\mu$ L of 0.25 M = 10 μmol) can be used in each coupling cycle of deoxy residues relative to polymer-bound 5'-hydroxyl. Average coupling yields on the 394 Applied Biosystems, Inc. synthesizer, determined by colorimetric quantitation of the trityl fractions, are typically 97.5-99%. Other oligonucleotide synthesis reagents for the 394 Applied Biosystems, Inc. synthesizer include the following: detritylation solution is 3% TCA in methylene chloride (ABI); capping is performed with 16% N-methyl imidazole in THF (ABI) and 10% acetic anhydride/10% 2,6-lutidine in THF (ABI); and oxidation solution is 16.9 mM I<sub>2</sub>, 49 mM pyridine, 9% water in THF (PERSEPTIVE™). Burdick & Jackson Synthesis Grade acetonitrile is used directly from the reagent bottle. S-Ethyltetrazole solution (0.25 M in acetonitrile) is made up from the solid obtained from American International Chemical, Inc. Alternately, for the introduction of phosphorothicate linkages, Beaucage reagent (3H-1,2-Benzodithiol-3-one 1,1-dioxide, 0.05 M in acetonitrile) is used.

Deprotection of the DNA-based oligonucleotides is performed as follows: the polymer-bound trityl-on oligoribonucleotide is transferred to a 4 mL glass screw top vial and suspended in a solution of 40% aq. methylamine (1 mL) at 65 °C for 10 min. After cooling to -20 °C, the supernatant is removed from the polymer support. The support is washed three times with 1.0 mL of EtOH:MeCN:H2O/3:1:1, vortexed and the supernatant is then added to the first supernatant. The combined supernatants, containing the oligoribonucleotide, are dried to a white powder.

The method of synthesis used for RNA including certain siNA molecules of the invention follows the procedure as described in Usman et al., 1987, J. Am. Chem. Soc., 109, 7845; Scaringe et al., 1990, Nucleic Acids Res., 18, 5433; and Wincott et al., 1995, Nucleic Acids Res. 23, 2677-2684 Wincott et al., 1997, Methods Mol. Bio., 74, 59, and makes use of common nucleic acid protecting and coupling groups, such as dimethoxytrityl at the 5'-end, and phosphoramidites at the 3'-end. In a non-limiting example, small scale syntheses are conducted on a 394 Applied Biosystems, Inc. synthesizer using a 0.2 µmol scale protocol with a 7.5 min coupling step for alkylsilyl

5

10

15

20

25

protected nucleotides and a 2.5 min coupling step for 2'-O-methylated nucleotides. Table II outlines the amounts and the contact times of the reagents used in the synthesis cycle. Alternatively, syntheses at the 0.2 µmol scale can be done on a 96-well plate synthesizer, such as the instrument produced by Protogene (Palo Alto, CA) with minimal modification A 33-fold excess (60  $\mu$ L of 0.11 M = 6.6  $\mu$ mol) of 2'-O-methyl to the cycle. phosphoramidite and a 75-fold excess of S-ethyl tetrazole (60  $\mu$ L of 0.25 M = 15  $\mu$ mol) can be used in each coupling cycle of 2'-O-methyl residues relative to polymer-bound 5'hydroxyl. A 66-fold excess (120  $\mu$ L of 0.11 M = 13.2  $\mu$ mol) of alkylsilyl (ribo) protected phosphoramidite and a 150-fold excess of S-ethyl tetrazole (120  $\mu$ L of 0.25 M = 30  $\mu$ mol) can be used in each coupling cycle of ribo residues relative to polymer-bound 5'hydroxyl. Average coupling yields on the 394 Applied Biosystems, Inc. synthesizer, determined by colorimetric quantitation of the trityl fractions, are typically 97.5-99%. Other oligonucleotide synthesis reagents for the 394 Applied Biosystems, Inc. synthesizer include the following: detritylation solution is 3% TCA in methylene chloride (ABI); capping is performed with 16% N-methyl imidazole in THF (ABI) and 10% acetic anhydride/10% 2,6-lutidine in THF (ABI); oxidation solution is 16.9 mM I2, 49 mM pyridine, 9% water in THF (PERSEPTIVE™). Burdick & Jackson Synthesis Grade acetonitrile is used directly from the reagent bottle. S-Ethyltetrazole solution (0.25 M in acetonitrile) is made up from the solid obtained from American International Chemical, Inc. Alternately, for the introduction of phosphorothioate linkages, Beaucage reagent (3H-1,2-Benzodithiol-3-one 1,1-dioxide0.05 M in acetonitrile) is used.

Deprotection of the RNA is performed using either a two-pot or one-pot protocol. For the two-pot protocol, the polymer-bound trityl-on oligoribonucleotide is transferred to a 4 mL glass screw top vial and suspended in a solution of 40% aq. methylamine (1 mL) at 65 °C for 10 min. After cooling to -20 °C, the supernatant is removed from the polymer support. The support is washed three times with 1.0 mL of EtOH:MeCN:H2O/3:1:1, vortexed and the supernatant is then added to the first supernatant. The combined supernatants, containing the oligoribonucleotide, are dried to a white powder. The base deprotected oligoribonucleotide is resuspended in anhydrous TEA/HF/NMP solution (300 μL of a solution of 1.5 mL N-methylpyrrolidinone, 750 μL TEA and 1 mL TEA•3HF to provide a 1.4 M HF concentration) and heated to 65 °C. After 1.5 h, the oligomer is quenched with 1.5 M NH<sub>4</sub>HCO<sub>3</sub>.

5

10

15

20

25

Alternatively, for the one-pot protocol, the polymer-bound trityl-on oligoribonucleotide is transferred to a 4 mL glass screw top vial and suspended in a solution of 33% ethanolic methylamine/DMSO: 1/1 (0.8 mL) at 65 °C for 15 min. The vial is brought to rt. TEA•3HF (0.1 mL) is added and the vial is heated at 65 °C for 15 min. The sample is cooled at -20 °C and then quenched with 1.5 M NH<sub>4</sub>HCO<sub>3</sub>.

For purification of the trityl-on oligomers, the quenched NH<sub>4</sub>HCO<sub>3</sub> solution is loaded onto a C-18 containing cartridge that had been prewashed with acetonitrile followed by 50 mM TEAA. After washing the loaded cartridge with water, the RNA is detritylated with 0.5% TFA for 13 min. The cartridge is then washed again with water, salt exchanged with 1 M NaCl and washed with water again. The oligonucleotide is then eluted with 30% acetonitrile.

The average stepwise coupling yields are typically >98% (Wincott et al., 1995 Nucleic Acids Res. 23, 2677-2684). Those of ordinary skill in the art will recognize that the scale of synthesis can be adapted to be larger or smaller than the example described above including but not limited to 96-well format.

Alternatively, the nucleic acid molecules of the present invention can be synthesized separately and joined together post-synthetically, for example, by ligation (Moore et al., 1992, Science 256, 9923; Draper et al., International PCT publication No. WO 93/23569; Shabarova et al., 1991, Nucleic Acids Research 19, 4247; Bellon et al., 1997, Nucleosides & Nucleotides, 16, 951; Bellon et al., 1997, Bioconjugate Chem. 8, 204), or by hybridization following synthesis and/or deprotection.

The siNA molecules of the invention can also be synthesized via a tandem synthesis methodology as described in Example 1 herein, wherein both siNA strands are synthesized as a single contiguous oligonucleotide fragment or strand separated by a cleavable linker which is subsequently cleaved to provide separate siNA fragments or strands that hybridize and permit purification of the siNA duplex. The linker can be a polynucleotide linker or a non-nucleotide linker. The tandem synthesis of siNA as described herein can be readily adapted to both multiwell/multiplate synthesis platforms such as 96 well or similarly larger multi-well platforms. The tandem synthesis of siNA as

5

10

15

20

described herein can also be readily adapted to large scale synthesis platforms employing batch reactors, synthesis columns and the like.

A siNA molecule can also be assembled from two distinct nucleic acid strands or fragments wherein one fragment includes the sense region and the second fragment includes the antisense region of the RNA molecule.

The nucleic acid molecules of the present invention can be modified extensively to enhance stability by modification with nuclease resistant groups, for example, 2'-amino, 2'-C-allyl, 2'-fluoro, 2'-O-methyl, 2'-H (for a review see Usman and Cedergren, 1992, TIBS 17, 34; Usman et al., 1994, Nucleic Acids Symp. Ser. 31, 163). siNA constructs can be purified by gel electrophoresis using general methods or can be purified by high pressure liquid chromatography (HPLC; see Wincott et al., supra, the totality of which is hereby incorporated herein by reference) and re-suspended in water.

In another aspect of the invention, siNA molecules of the invention are expressed from transcription units inserted into DNA or RNA vectors. The recombinant vectors can be DNA plasmids or viral vectors. siNA expressing viral vectors can be constructed based on, but not limited to, adeno-associated virus, retrovirus, adenovirus, or alphavirus. The recombinant vectors capable of expressing the siNA molecules can be delivered as described herein, and persist in target cells. Alternatively, viral vectors can be used that provide for transient expression of siNA molecules.

## 20 Optimizing Activity of the nucleic acid molecule of the invention.

Chemically synthesizing nucleic acid molecules with modifications (base, sugar and/or phosphate) can prevent their degradation by serum ribonucleases, which can increase their potency (see e.g., Eckstein et al., International Publication No. WO 92/07065; Perrault et al., 1990 Nature 344, 565; Pieken et al., 1991, Science 253, 314; Usman and Cedergren, 1992, Trends in Biochem. Sci. 17, 334; Usman et al., International Publication No. WO 93/15187; and Rossi et al., International Publication No. WO 91/03162; Sproat, U.S. Pat. No. 5,334,711; Gold et al., U.S. Pat. No. 6,300,074; and Burgin et al., supra; all of which are incorporated by reference herein). All of the above references describe various chemical modifications that can be made to the base, phosphate and/or sugar moieties of the nucleic acid molecules described herein.

5

10

15

25

Modifications that enhance their efficacy in cells, and removal of bases from nucleic acid molecules to shorten oligonucleotide synthesis times and reduce chemical requirements are desired.

There are several examples in the art describing sugar, base and phosphate modifications that can be introduced into nucleic acid molecules with significant 5 enhancement in their nuclease stability and efficacy. For example, oligonucleotides are modified to enhance stability and/or enhance biological activity by modification with nuclease resistant groups, for example, 2'-amino, 2'-C-allyl, 2'-fluoro, 2'-O-methyl, 2'-Oallyl, 2'-H, nucleotide base modifications (for a review see Usman and Cedergren, 1992, TIBS. 17, 34; Usman et al., 1994, Nucleic Acids Symp. Ser. 31, 163; Burgin et al., 1996, 10 Biochemistry, 35, 14090). Sugar modification of nucleic acid molecules have been extensively described in the art (see Eckstein et al., International Publication PCT No. WO 92/07065; Perrault et al. Nature, 1990, 344, 565-568; Pieken et al. Science, 1991, 253, 314-317; Usman and Cedergren, Trends in Biochem. Sci., 1992, 17, 334-339; Usman et al. International Publication PCT No. WO 93/15187; Sproat, U.S. Pat. No. 15 5,334,711 and Beigelman et al., 1995, J. Biol. Chem., 270, 25702; Beigelman et al., International PCT publication No. WO 97/26270; Beigelman et al., U.S. Pat. No. 5,716,824; Usman et al., U.S. Pat. No. 5,627,053; Woolf et al., International PCT Publication No. WO 98/13526; Thompson et al., USSN 60/082,404 which was filed on April 20, 1998; Karpeisky et al., 1998, Tetrahedron Lett., 39, 1131; Earnshaw and Gait, 20 1998, Biopolymers (Nucleic Acid Sciences), 48, 39-55; Verma and Eckstein, 1998, Annu. Rev. Biochem., 67, 99-134; and Burlina et al., 1997, Bioorg. Med. Chem., 5, 1999-2010; all of the references are hereby incorporated in their totality by reference herein). Such publications describe general methods and strategies to determine the location of incorporation of sugar, base and/or phosphate modifications and the like into nucleic acid 25 molecules without modulating catalysis, and are incorporated by reference herein. In view of such teachings, similar modifications can be used as described herein to modify the siNA nucleic acid molecules of the instant invention so long as the ability of siNA to promote RNAi is cells is not significantly inhibited.

While chemical modification of oligonucleotide internucleotide linkages with phosphorothioate, phosphorodithioate, and/or 5'-methylphosphonate linkages improves stability, excessive modifications can cause some toxicity or decreased activity.

Therefore, when designing nucleic acid molecules, the amount of these internucleotide linkages should be minimized. The reduction in the concentration of these linkages should lower toxicity, resulting in increased efficacy and higher specificity of these molecules.

Short interfering nucleic acid (siNA) molecules having chemical modifications that maintain or enhance activity are provided. Such a nucleic acid is also generally more resistant to nucleases than an unmodified nucleic acid. Accordingly, the *in vitro* and/or *in vivo* activity should not be significantly lowered. In cases in which modulation is the goal, therapeutic nucleic acid molecules delivered exogenously should optimally be stable within cells until translation of the target RNA has been modulated long enough to reduce the levels of the undesirable protein. This period of time varies between hours to days depending upon the disease state. Improvements in the chemical synthesis of RNA and DNA (Wincott *et al.*, 1995, *Nucleic Acids Res.* 23, 2677; Caruthers *et al.*, 1992, *Methods in Enzymology* 211,3-19 (incorporated by reference herein)) have expanded the ability to modify nucleic acid molecules by introducing nucleotide modifications to enhance their nuclease stability, as described above.

In one embodiment, nucleic acid molecules of the invention include one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) G-clamp nucleotides. A G-clamp nucleotide is a modified cytosine analog wherein the modifications confer the ability to hydrogen bond both Watson-Crick and Hoogsteen faces of a complementary guanine within a duplex, see for example Lin and Matteucci, 1998, J. Am. Chem. Soc., 120, 8531-8532. A single G-clamp analog substitution within an oligonucleotide can result in substantially enhanced helical thermal stability and mismatch discrimination when hybridized to complementary oligonucleotides. The inclusion of such nucleotides in nucleic acid molecules of the invention results in both enhanced affinity and specificity to nucleic acid targets, complementary sequences, or template strands. In another embodiment, nucleic acid molecules of the invention include one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) LNA "locked nucleic acid" nucleotides such as a 2', 4'-C methylene bicyclo nucleotide (see for example Wengel et al., International PCT Publication No. WO 00/66604 and WO 99/14226).

5

10

15

20

25

In another embodiment, the invention features conjugates and/or complexes of siNA molecules of the invention. Such conjugates and/or complexes can be used to facilitate delivery of siNA molecules into a biological system, such as a cell. conjugates and complexes provided by the instant invention can impart therapeutic activity by transferring therapeutic compounds across cellular membranes, altering the pharmacokinetics, and/or modulating the localization of nucleic acid molecules of the The present invention encompasses the design and synthesis of novel invention. conjugates and complexes for the delivery of molecules, including, but not limited to, small molecules, lipids, phospholipids, nucleosides, nucleotides, nucleic acids, antibodies, toxins, negatively charged polymers and other polymers, for example proteins, peptides, hormones, carbohydrates, polyethylene glycols, or polyamines, across cellular In general, the transporters described are designed to be used either membranes. individually or as part of a multi-component system, with or without degradable linkers. These compounds are expected to improve delivery and/or localization of nucleic acid molecules of the invention into a number of cell types originating from different tissues, in the presence or absence of serum (see Sullenger and Cech, U.S. Pat. No. 5,854,038). Conjugates of the molecules described herein can be attached to biologically active molecules via linkers that are biodegradable, such as biodegradable nucleic acid linker molecules.

The term "biodegradable linker" as used herein, refers to a nucleic acid or non-nucleic acid linker molecule that is designed as a biodegradable linker to connect one molecule to another molecule, for example, a biologically active molecule to a siNA molecule of the invention or the sense and antisense strands of a siNA molecule of the invention. The biodegradable linker is designed such that its stability can be modulated for a particular purpose, such as delivery to a particular tissue or cell type. The stability of a nucleic acid-based biodegradable linker molecule can be modulated by using various chemistries, for example combinations of ribonucleotides, deoxyribonucleotides, and chemically-modified nucleotides, such as 2'-O-methyl, 2'-fluoro, 2'-amino, 2'-O-amino, 2'-C-allyl, 2'-O-allyl, and other 2'-modified or base modified nucleotides. The biodegradable nucleic acid linker molecule can be a dimer, trimer, tetramer or longer nucleic acid molecule, for example, an oligonucleotide of about 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, or 20 nucleotides in length, or can comprise a single

5

10

15

20

25

nucleotide with a phosphorus-based linkage, for example, a phosphoramidate or phosphodiester linkage. The biodegradable nucleic acid linker molecule can also comprise nucleic acid backbone, nucleic acid sugar, or nucleic acid base modifications.

The term "biodegradable" as used herein, refers to degradation in a biological system, for example enzymatic degradation or chemical degradation.

The term "biologically active molecule" as used herein, refers to compounds or molecules that are capable of eliciting or modifying a biological response in a system. Non-limiting examples of biologically active siNA molecules either alone or in combination with other molecules contemplated by the instant invention include therapeutically active molecules such as antibodies, hormones, antivirals, peptides, proteins, chemotherapeutics, small molecules, vitamins, co-factors, nucleosides, nucleotides, oligonucleotides, enzymatic nucleic acids, antisense nucleic acids, triplex forming oligonucleotides, 2,5-A chimeras, siNA, dsRNA, allozymes, aptamers, decoys and analogs thereof. Biologically active molecules of the invention also include molecules capable of modulating the pharmacokinetics and/or pharmacodynamics of other biologically active molecules, for example, lipids and polymers such as polyamines, polyamides, polyethylene glycol and other polyethers.

\*

The term "phospholipid" as used herein, refers to a hydrophobic molecule comprising at least one phosphorus group. For example, a phospholipid can comprise a phosphorus-containing group and saturated or unsaturated alkyl group, optionally substituted with OH, COOH, oxo, amine, or substituted or unsubstituted aryl groups.

Therapeutic nucleic acid molecules (e.g., siNA molecules) delivered exogenously optimally are stable within cells until reverse transcription of the RNA has been modulated long enough to reduce the levels of the RNA transcript. The nucleic acid molecules are resistant to nucleases in order to function as effective intracellular therapeutic agents. Improvements in the chemical synthesis of nucleic acid molecules described in the instant invention and in the art have expanded the ability to modify nucleic acid molecules by introducing nucleotide modifications to enhance their nuclease stability as described above.

5

10

15

20

In yet another embodiment, siNA molecules having chemical modifications that maintain or enhance enzymatic activity of proteins involved in RNAi are provided. Such nucleic acids are also generally more resistant to nucleases than unmodified nucleic acids. Thus, *in vitro* and/or *in vivo* the activity should not be significantly lowered.

Use of the nucleic acid-based molecules of the invention will lead to better treatment of the disease progression by affording the possibility of combination therapies (e.g., multiple siNA molecules targeted to different genes; nucleic acid molecules coupled with known small molecule modulators; or intermittent treatment with combinations of molecules, including different motifs and/or other chemical or biological molecules). The treatment of subjects with siNA molecules can also include combinations of different types of nucleic acid molecules, such as enzymatic nucleic acid molecules (ribozymes), allozymes, antisense, 2,5-A oligoadenylate, decoys, and aptamers.

In another aspect a siNA molecule of the invention comprises one or more 5' and/or a 3'- cap structure, for example on only the sense siNA strand, the antisense siNA strand, or both siNA strands.

By "cap structure" is meant chemical modifications, which have been incorporated at either terminus of the oligonucleotide (see, for example, Adamic et al., U.S. Pat. No. 5,998,203, incorporated by reference herein). These terminal modifications protect the nucleic acid molecule from exonuclease degradation, and may help in delivery and/or localization within a cell. The cap may be present at the 5'-terminus (5'-cap) or at the 3'terminal (3'-cap) or may be present on both termini. In non-limiting examples, the 5'-cap is selected from the group consisting of glyceryl, inverted deoxy abasic residue (moiety); 4',5'-methylene nucleotide; 1-(beta-D-erythrofuranosyl) nucleotide, 4'-thio nucleotide; carbocyclic nucleotide; 1,5-anhydrohexitol nucleotide; L-nucleotides; alpha-nucleotides; modified base nucleotide; phosphorodithioate linkage; threo-pentofuranosyl nucleotide; acyclic 3',4'-seco nucleotide; acyclic 3,4-dihydroxybutyl nucleotide; acyclic 3,5dihydroxypentyl nucleotide, 3'-3'-inverted nucleotide moiety; 3'-3'-inverted abasic moiety; 3'-2'-inverted nucleotide moiety; 3'-2'-inverted abasic moiety; 1,4-butanediol phosphate; 3'-phosphoramidate; hexylphosphate; aminohexyl phosphate; 3'-phosphate; 3'phosphorothioate; phosphorodithioate; or bridging or non-bridging methylphosphonate moiety.

5

10

15

20

25

In non-limiting examples, the 3'-cap is selected from the group consisting of glyceryl, inverted deoxy abasic residue (moiety), 4',5'-methylene nucleotide; 1-(beta-D-erythrofuranosyl) nucleotide; 4'-thio nucleotide, carbocyclic nucleotide; 5'-amino-alkyl phosphate; 1,3-diamino-2-propyl phosphate; 3-aminopropyl phosphate; 6-aminohexyl phosphate; 1,2-aminododecyl phosphate; hydroxypropyl phosphate; 1,5-anhydrohexitol nucleotide; L-nucleotide; alpha-nucleotide; modified base nucleotide; phosphorodithioate; threo-pentofuranosyl nucleotide; acyclic 3',4'-seco nucleotide; 3,4-dihydroxybutyl nucleotide; 3,5-dihydroxypentyl nucleotide, 5'-5'-inverted nucleotide moiety; 5'-5'-inverted abasic moiety; 5'-phosphoramidate; 5'-phosphorothioate; 1,4-butanediol phosphate; 5'-amino; bridging and/or non-bridging 5'-phosphoramidate, phosphorothioate and/or phosphorodithioate, bridging or non bridging methylphosphonate and 5'-mercapto moieties (for more details see Beaucage and Iyer, 1993, Tetrahedron 49, 1925; incorporated by reference herein).

By the term "non-nucleotide" is meant any group or compound which can be incorporated into a nucleic acid chain in the place of one or more nucleotide units, including either sugar and/or phosphate substitutions, and allows the remaining bases to exhibit their enzymatic activity. The group or compound is abasic in that it does not contain a commonly recognized nucleotide base, such as adenosine, guanine, cytosine, uracil or thymine and therefore lacks a base at the 1'-position.

An "alkyl" group refers to a saturated aliphatic hydrocarbon, including straight-chain, branched-chain, and cyclic alkyl groups. Preferably, the alkyl group has 1 to 12 carbons. More preferably, it is a lower alkyl of from 1 to 7 carbons, more preferably 1 to 4 carbons. The alkyl group can be substituted or unsubstituted. When substituted the substituted group(s) is preferably, hydroxyl, cyano, alkoxy, =0, =S, NO2 or N(CH3)2, amino, or SH. The term also includes alkenyl groups that are unsaturated hydrocarbon groups containing at least one carbon-carbon double bond, including straight-chain, branched-chain, and cyclic groups. Preferably, the alkenyl group has 1 to 12 carbons. More preferably, it is a lower alkenyl of from 1 to 7 carbons, more preferably 1 to 4 carbons. The alkenyl group may be substituted or unsubstituted. When substituted the substituted group(s) is preferably, hydroxyl, cyano, alkoxy, =0, =S, NO2, halogen, N(CH3)2, amino, or SH. The term "alkyl" also includes alkynyl groups that have an

5

10

15

20

25

unsaturated hydrocarbon group containing at least one carbon-carbon triple bond, including straight-chain, branched-chain, and cyclic groups. Preferably, the alkynyl group has 1 to 12 carbons. More preferably, it is a lower alkynyl of from 1 to 7 carbons, more preferably 1 to 4 carbons. The alkynyl group may be substituted or unsubstituted. When substituted the substituted group(s) is preferably, hydroxyl, cyano, alkoxy, =O, =S, NO2 or N(CH3)2, amino or SH.

Such alkyl groups can also include aryl, alkylaryl, carbocyclic aryl, heterocyclic aryl, amide and ester groups. An "aryl" group refers to an aromatic group that has at least one ring having a conjugated pi electron system and includes carbocyclic aryl, heterocyclic aryl and biaryl groups, all of which may be optionally substituted. The preferred substituent(s) of aryl groups are halogen, trihalomethyl, hydroxyl, SH, OH, cyano, alkoxy, alkyl, alkenyl, alkynyl, and amino groups. An "alkylaryl" group refers to an alkyl group (as described above) covalently joined to an aryl group (as described above). Carbocyclic aryl groups are groups wherein the ring atoms on the aromatic ring are all carbon atoms. The carbon atoms are optionally substituted. Heterocyclic aryl groups are groups having from 1 to 3 heteroatoms as ring atoms in the aromatic ring and the remainder of the ring atoms are carbon atoms. Suitable heteroatoms include oxygen, sulfur, and nitrogen, and include furanyl, thienyl, pyridyl, pyrrolyl, N-lower alkyl pyrrolo, pyrimidyl, pyrazinyl, imidazolyl and the like, all optionally substituted. An "amide" refers to an -C(O)-NH-R, where R is either alkyl, aryl, alkylaryl or hydrogen.

By "nucleotide" as used herein is as recognized in the art to include natural bases (standard), and modified bases well known in the art. Such bases are generally located at the 1' position of a nucleotide sugar moiety. Nucleotides generally comprise a base, sugar and a phosphate group. The nucleotides can be unmodified or modified at the sugar, phosphate and/or base moiety, (also referred to interchangeably as nucleotide analogs, modified nucleotides, non-natural nucleotides, non-standard nucleotides and other; see, for example, Usman and McSwiggen, *supra*; Eckstein *et al.*, International PCT Publication No. WO 92/07065; Usman *et al.*, International PCT Publication No. WO 93/15187; Uhlman & Peyman, *supra*, all are hereby incorporated by reference herein). There are several examples of modified nucleic acid bases known in the art as summarized by Limbach *et al.*, 1994, *Nucleic Acids Res.* 22, 2183. Some of the non-

5

10

15

20

25

limiting examples of base modifications that can be introduced into nucleic acid molecules include, inosine, purine, pyridin-4-one, pyridin-2-one, phenyl, pseudouracil, 2, 4, 6-trimethoxy benzene, 3-methyl uracil, dihydrouridine, naphthyl, aminophenyl, 5-alkylcytidines (e.g., 5-methylcytidine), 5-alkyluridines (e.g., ribothymidine), 5-halouridine (e.g., 5-bromouridine) or 6-azapyrimidines or 6-alkylpyrimidines (e.g. 6-methyluridine), propyne, and others (Burgin et al., 1996, Biochemistry, 35, 14090; Uhlman & Peyman, supra). By "modified bases" in this aspect is meant nucleotide bases other than adenine, guanine, cytosine and uracil at 1' position or their equivalents.

In one embodiment, the invention features modified siNA molecules, with phosphate backbone modifications comprising one or more phosphorothioate, phosphorodithioate, methylphosphonate, phosphotriester, morpholino, amidate carbamate, carboxymethyl, acetamidate, polyamide, sulfonate, sulfonamide, sulfamate, formacetal, thioformacetal, and/or alkylsilyl, substitutions. For a review of oligonucleotide backbone modifications, see Hunziker and Leumann, 1995, *Nucleic Acid Analogues: Synthesis and Properties*, in *Modern Synthetic Methods*, VCH, 331-417, and Mesmaeker et al., 1994, *Novel Backbone Replacements for Oligonucleotides*, in *Carbohydrate Modifications in Antisense Research*, ACS, 24-39.

By "abasic" is meant sugar moieties lacking a base or having other chemical groups in place of a base at the 1' position, see for example Adamic *et al.*, U.S. Pat. No. 5,998,203.

By "unmodified nucleoside" is meant one of the bases adenine, cytosine, guanine, thymine, or uracil joined to the 1' carbon of  $\beta$ -D-ribo-furanose.

By "modified nucleoside" is meant any nucleotide base which contains a modification in the chemical structure of an unmodified nucleotide base, sugar and/or phosphate. Non-limiting examples of modified nucleotides are shown by Formulae I-VII and/or other modifications described herein.

In connection with 2'-modified nucleotides as described for the present invention, by "amino" is meant 2'-NH<sub>2</sub> or 2'-O- NH<sub>2</sub>, which can be modified or unmodified. Such modified groups are described, for example, in Eckstein *et al.*, U.S. Pat. No. 5,672,695

5

10

15

20

and Matulic-Adamic et al., U.S. Pat. No. 6,248,878, which are both incorporated by reference in their entireties.

Various modifications to nucleic acid siNA structure can be made to enhance the utility of these molecules. Such modifications will enhance shelf-life, half-life *in vitro*, stability, and ease of introduction of such oligonucleotides to the target site, *e.g.*, to enhance penetration of cellular membranes, and confer the ability to recognize and bind to targeted cells.

## Administration of Nucleic Acid Molecules

A siNA molecule of the invention can be adapted for use to treat any disease, infection or condition associated with gene expression, and other indications that can respond to the level of gene product in a cell or tissue, alone or in combination with other therapies. For example, a siNA molecule can comprise a delivery vehicle, including liposomes, for administration to a subject, carriers and diluents and their salts, and/or can be present in pharmaceutically acceptable formulations. Methods for the delivery of nucleic acid molecules are described in Akhtar et al., 1992, Trends Cell Bio., 2, 139; Delivery Strategies for Antisense Oligonucleotide Therapeutics, ed. Akhtar, 1995, Maurer et al., 1999, Mol. Membr. Biol., 16, 129-140; Hofland and Huang, 1999, Handb. Exp. Pharmacol., 137, 165-192; and Lee et al., 2000, ACS Symp. Ser., 752, 184-192, all of which are incorporated herein by reference. Beigelman et al., U.S. Pat. No. 6,395,713 and Sullivan et al., PCT WO 94/02595 further describe the general methods for delivery of nucleic acid molecules. These protocols can be utilized for the delivery of virtually any nucleic acid molecule. Nucleic acid molecules can be administered to cells by a variety of methods known to those of skill in the art, including, but not restricted to, encapsulation in liposomes, by iontophoresis, or by incorporation into other vehicles, such as hydrogels, cyclodextrins (see for example Gonzalez et al., 1999, Bioconjugate Chem., 10, 1068-1074), biodegradable nanocapsules, and bioadhesive microspheres, or by proteinaceous vectors (O'Hare and Normand, International PCT Publication No. WO 00/53722). Alternatively, the nucleic acid/vehicle combination is locally delivered by direct injection or by use of an infusion pump. Direct injection of the nucleic acid molecules of the invention, whether subcutaneous, intramuscular, or intradermal, can take place using standard needle and syringe methodologies, or by needle-free technologies

5

10

15

20

25

such as those described in Conry et al., 1999, Clin. Cancer Res., 5, 2330-2337 and Barry et al., International PCT Publication No. WO 99/31262. Many examples in the art describe CNS delivery methods of oligonucleotides by osmotic pump, (see Chun et al., 1998, Neuroscience Letters, 257, 135-138, D'Aldin et al., 1998, Mol. Brain Research, 55, 151-164, Dryden et al., 1998, J. Endocrinol., 157, 169-175, Ghirnikar et al., 1998, Neuroscience Letters, 247, 21-24) or direct infusion (Broaddus et al., 1997, Neurosurg. Focus, 3, article 4). Other routes of delivery include, but are not limited to oral (tablet or pill form) and/or intrathecal delivery (Gold, 1997, Neuroscience, 76, 1153-1158). More detailed descriptions of nucleic acid delivery and administration are provided in Sullivan et al., supra, Draper et al., PCT WO93/23569, Beigelman et al., PCT WO99/05094, and Klimuk et al., PCT WO99/04819 all of which have been incorporated by reference herein. The molecules of the instant invention can be used as pharmaceutical agents. Pharmaceutical agents prevent, modulate the occurrence, or treat (alleviate a symptom to some extent, preferably all of the symptoms) of a disease state in a subject.

In addition, the invention features the use of methods to deliver the nucleic acid molecules of the instant invention to hematopoietic cells, including monocytes and lymphocytes. These methods are described in detail by Hartmann *et al.*, 1998, *J. Phamacol. Exp. Ther.*, 285(2), 920-928; Kronenwett *et al.*, 1998, *Blood*, 91(3), 852-862; Filion and Phillips, 1997, *Biochim. Biophys. Acta.*, 1329(2), 345-356; Ma and Wei, 1996, *Leuk. Res.*, 20(11/12), 925-930; and Bongartz *et al.*, 1994, *Nucleic Acids Research*, 22(22), 4681-8. Such methods, as described above, include the use of free oligonucleitide, cationic lipid formulations, liposome formulations including pH sensitive liposomes and immunoliposomes, and bioconjugates including oligonucleotides conjugated to fusogenic peptides, for the transfection of hematopoietic cells with oligonucleotides.

Thus, the invention features a pharmaceutical composition comprising one or more nucleic acid(s) of the invention in an acceptable carrier, such as a stabilizer, buffer, and the like. The polynucleotides of the invention can be administered (e.g., RNA, DNA or protein) and introduced into a subject by any standard means, with or without stabilizers, buffers, and the like, to form a pharmaceutical composition. When it is desired to use a liposome delivery mechanism, standard protocols for formation of liposomes can be followed. The compositions of the present invention can also be formulated and used as

5

10

15

20

25

tablets, capsules or elixirs for oral administration, suppositories for rectal administration, sterile solutions, suspensions for injectable administration, and the other compositions known in the art.

The present invention also includes pharmaceutically acceptable formulations of the compounds described. These formulations include salts of the above compounds, *e.g.*, acid addition salts, for example, salts of hydrochloric, hydrobromic, acetic acid, and benzene sulfonic acid.

A pharmacological composition or formulation refers to a composition or formulation in a form suitable for administration, e.g., systemic administration, into a cell or subject, including for example a human. Suitable forms, in part, depend upon the use or the route of entry, for example oral, transdermal, or by injection. Such forms should not prevent the composition or formulation from reaching a target cell (i.e., a cell to which the negatively charged nucleic acid is desirable for delivery). For example, pharmacological compositions injected into the blood stream should be soluble. Other factors are known in the art, and include considerations such as toxicity and forms that prevent the composition or formulation from exerting its effect.

By "systemic administration" is meant *in vivo* systemic absorption or accumulation of drugs in the blood stream followed by distribution throughout the entire body. Administration routes that lead to systemic absorption include, without limitation: intravenous, subcutaneous, intraperitoneal, inhalation, oral, intrapulmonary and intramuscular. Each of these administration routes exposes the siNA molecules of the invention to an accessible diseased tissue. The rate of entry of a drug into the circulation has been shown to be a function of molecular weight or size. The use of a liposome or other drug carrier comprising the compounds of the instant invention can potentially localize the drug, for example, in certain tissue types, such as the tissues of the reticular endothelial system (RES). A liposome formulation that can facilitate the association of drug with the surface of cells, such as, lymphocytes and macrophages is also useful. This approach can provide enhanced delivery of the drug to target cells by taking advantage of the specificity of macrophage and lymphocyte immune recognition of abnormal cells, such as cells producing excess MDR.

5

10

15

20

25

"pharmaceutically acceptable formulation" is meant, a composition or formulation that allows for the effective distribution of the nucleic acid molecules of the instant invention in the physical location most suitable for their desired activity. Nonlimiting examples of agents suitable for formulation with the nucleic acid molecules of the instant invention include: P-glycoprotein inhibitors (such as Pluronic P85), which can enhance entry of drugs into the CNS (Jolliet-Riant and Tillement, 1999, Fundam. Clin. Pharmacol., 13, 16-26); biodegradable polymers, such as poly (DL-lactide-coglycolide) microspheres for sustained release delivery after intracerebral implantation (Emerich, DF et al, 1999, Cell Transplant, 8, 47-58) (Alkermes, Inc. Cambridge, MA); and loaded nanoparticles, such as those made of polybutylcyanoacrylate, which can deliver drugs across the blood brain barrier and can alter neuronal uptake mechanisms (Prog Neuropsychopharmacol Biol Psychiatry, 23, 941-949, 1999). Other non-limiting examples of delivery strategies for the nucleic acid molecules of the instant invention include material described in Boado et al., 1998, J. Pharm. Sci., 87, 1308-1315; Tyler et al., 1999, FEBS Lett., 421, 280-284; Pardridge et al., 1995, PNAS USA., 92, 5592-5596; Boado, 1995, Adv. Drug Delivery Rev., 15, 73-107; Aldrian-Herrada et al., 1998, Nucleic Acids Res., 26, 4910-4916; and Tyler et al., 1999, PNAS USA., 96, 7053-7058.

The invention also features the use of the composition comprising surface-modified liposomes containing poly (ethylene glycol) lipids (PEG-modified, or long-circulating liposomes or stealth liposomes). These formulations offer a method for increasing the accumulation of drugs in target tissues. This class of drug carriers resists opsonization and elimination by the mononuclear phagocytic system (MPS or RES), thereby enabling longer blood circulation times and enhanced tissue exposure for the encapsulated drug (Lasic et al. Chem. Rev. 1995, 95, 2601-2627; Ishiwata et al., Chem. Pharm. Bull. 1995, 43, 1005-1011). Such liposomes have been shown to accumulate selectively in tumors, presumably by extravasation and capture in the neovascularized target tissues (Lasic et al., Science 1995, 267, 1275-1276; Oku et al., 1995, Biochim. Biophys. Acta, 1238, 86-90). The long-circulating liposomes the enhance pharmacokinetics and pharmacodynamics of DNA and RNA, particularly compared to conventional cationic liposomes which are known to accumulate in tissues of the MPS (Liu et al., J. Biol. Chem. 1995, 42, 24864-24870; Choi et al., International PCT Publication No. WO 96/10391; Ansell et al., International PCT Publication No. WO 96/10390; Holland et al.,

5

10

15

20

25

International PCT Publication No. WO 96/10392). Long-circulating liposomes are also likely to protect drugs from nuclease degradation to a greater extent compared to cationic liposomes, based on their ability to avoid accumulation in metabolically aggressive MPS tissues such as the liver and spleen.

The present invention also includes compositions prepared for storage or administration that include a pharmaceutically effective amount of the desired compounds in a pharmaceutically acceptable carrier or diluent. Acceptable carriers or diluents for therapeutic use are well known in the pharmaceutical art, and are described, for example, in *Remington's Pharmaceutical Sciences*, Mack Publishing Co. (A.R. Gennaro edit. 1985), hereby incorporated by reference herein. For example, preservatives, stabilizers, dyes and flavoring agents can be provided. These include sodium benzoate, sorbic acid and esters of *p*-hydroxybenzoic acid. In addition, antioxidants and suspending agents can be used.

A pharmaceutically effective dose is that dose required to prevent, inhibit the occurrence, or treat (alleviate a symptom to some extent, preferably all of the symptoms) of a disease state. The pharmaceutically effective dose depends on the type of disease, the composition used, the route of administration, the type of mammal being treated, the physical characteristics of the specific mammal under consideration, concurrent medication, and other factors that those skilled in the medical arts will recognize. Generally, an amount between 0.1 mg/kg and 100 mg/kg body weight/day of active ingredients is administered dependent upon potency of the negatively charged polymer.

The nucleic acid molecules of the invention and formulations thereof can be administered orally, topically, parenterally, by inhalation or spray, or rectally in dosage unit formulations containing conventional non-toxic pharmaceutically acceptable carriers, adjuvants and/or vehicles. The term parenteral as used herein includes percutaneous, subcutaneous, intravascular (e.g., intravenous), intramuscular, or intrathecal injection or infusion techniques and the like. In addition, there is provided a pharmaceutical formulation comprising a nucleic acid molecule of the invention and a pharmaceutically acceptable carrier. One or more nucleic acid molecules of the invention can be present in association with one or more non-toxic pharmaceutically acceptable carriers and/or diluents and/or adjuvants, and if desired other active ingredients. The pharmaceutical

5

10

15

20

25

compositions containing nucleic acid molecules of the invention can be in a form suitable for oral use, for example, as tablets, troches, lozenges, aqueous or oily suspensions, dispersible powders or granules, emulsion, hard or soft capsules, or syrups or elixirs.

Compositions intended for oral use can be prepared according to any method known to the art for the manufacture of pharmaceutical compositions and such compositions can contain one or more such sweetening agents, flavoring agents, coloring agents or preservative agents in order to provide pharmaceutically elegant and palatable Tablets contain the active ingredient in admixture with non-toxic preparations. pharmaceutically acceptable excipients that are suitable for the manufacture of tablets. These excipients can be, for example, inert diluents; such as calcium carbonate, sodium carbonate, lactose, calcium phosphate or sodium phosphate; granulating and disintegrating agents, for example, corn starch, or alginic acid; binding agents, for example starch, gelatin or acacia; and lubricating agents, for example magnesium stearate, stearic acid or talc. The tablets can be uncoated or they can be coated by known techniques. In some cases such coatings can be prepared by known techniques to delay disintegration and absorption in the gastrointestinal tract and thereby provide a sustained action over a longer period. For example, a time delay material such as glyceryl monosterate or glyceryl distearate can be employed.

Formulations for oral use can also be presented as hard gelatin capsules wherein the active ingredient is mixed with an inert solid diluent, for example, calcium carbonate, calcium phosphate or kaolin, or as soft gelatin capsules wherein the active ingredient is mixed with water or an oil medium, for example peanut oil, liquid paraffin or olive oil.

Aqueous suspensions contain the active materials in a mixture with excipients suitable for the manufacture of aqueous suspensions. Such excipients are suspending agents, for example sodium carboxymethylcellulose, methylcellulose, hydropropylmethylcellulose, sodium alginate, polyvinylpyrrolidone, gum tragacanth and gum acacia; dispersing or wetting agents can be a naturally-occurring phosphatide, for example, lecithin, or condensation products of an alkylene oxide with fatty acids, for example polyoxyethylene stearate, or condensation products of ethylene oxide with long chain aliphatic alcohols, for example heptadecaethyleneoxycetanol, or condensation products of ethylene oxide with partial esters derived from fatty acids and a hexitol such as

5

10

15

20

25

polyoxyethylene sorbitol monooleate, or condensation products of ethylene oxide with partial esters derived from fatty acids and hexitol anhydrides, for example polyethylene sorbitan monooleate. The aqueous suspensions can also contain one or more preservatives, for example ethyl, or n-propyl p-hydroxybenzoate, one or more coloring agents, one or more flavoring agents, and one or more sweetening agents, such as sucrose or saccharin.

Oily suspensions can be formulated by suspending the active ingredients in a vegetable oil, for example arachis oil, olive oil, sesame oil or coconut oil, or in a mineral oil such as liquid paraffin. The oily suspensions can contain a thickening agent, for example beeswax, hard paraffin or cetyl alcohol. Sweetening agents and flavoring agents can be added to provide palatable oral preparations. These compositions can be preserved by the addition of an anti-oxidant such as ascorbic acid

Dispersible powders and granules suitable for preparation of an aqueous suspension by the addition of water provide the active ingredient in admixture with a dispersing or wetting agent, suspending agent and one or more preservatives. Suitable dispersing or wetting agents or suspending agents are exemplified by those already mentioned above. Additional excipients, for example sweetening, flavoring and coloring agents, can also be present.

Pharmaceutical compositions of the invention can also be in the form of oil-in-water emulsions. The oily phase can be a vegetable oil or a mineral oil or mixtures of these. Suitable emulsifying agents can be naturally-occurring gums, for example gum acacia or gum tragacanth, naturally-occurring phosphatides, for example soy bean, lecithin, and esters or partial esters derived from fatty acids and hexitol, anhydrides, for example sorbitan monooleate, and condensation products of the said partial esters with ethylene oxide, for example polyoxyethylene sorbitan monooleate. The emulsions can also contain sweetening and flavoring agents.

Syrups and elixirs can be formulated with sweetening agents, for example glycerol, propylene glycol, sorbitol, glucose or sucrose. Such formulations can also contain a demulcent, a preservative and flavoring and coloring agents. The pharmaceutical compositions can be in the form of a sterile injectable aqueous or oleaginous suspension. This suspension can be formulated according to the known art using those suitable

5

10

15

20

25

dispersing or wetting agents and suspending agents that have been mentioned above. The sterile injectable preparation can also be a sterile injectable solution or suspension in a non-toxic parentally acceptable diluent or solvent, for example as a solution in 1,3-butanediol. Among the acceptable vehicles and solvents that can be employed are water, Ringer's solution and isotonic sodium chloride solution. In addition, sterile, fixed oils are conventionally employed as a solvent or suspending medium. For this purpose, any bland fixed oil can be employed including synthetic mono-or diglycerides. In addition, fatty acids such as oleic acid find use in the preparation of injectables.

The nucleic acid molecules of the invention can also be administered in the form of suppositories, e.g., for rectal administration of the drug. These compositions can be prepared by mixing the drug with a suitable non-irritating excipient that is solid at ordinary temperatures but liquid at the rectal temperature and will therefore melt in the rectum to release the drug. Such materials include cocoa butter and polyethylene glycols.

Nucleic acid molecules of the invention can be administered parenterally in a sterile medium. The drug, depending on the vehicle and concentration used, can either be suspended or dissolved in the vehicle. Advantageously, adjuvants such as local anesthetics, preservatives and buffering agents can be dissolved in the vehicle.

Dosage levels of the order of from about 0.1 mg to about 140 mg per kilogram of body weight per day are useful in the treatment of the above-indicated conditions (about 0.5 mg to about 7 g per subject per day). The amount of active ingredient that can be combined with the carrier materials to produce a single dosage form varies depending upon the host treated and the particular mode of administration. Dosage unit forms generally contain between from about 1 mg to about 500 mg of an active ingredient.

It is understood that the specific dose level for any particular subject depends upon a variety of factors including the activity of the specific compound employed, the age, body weight, general health, sex, diet, time of administration, route of administration, and rate of excretion, drug combination and the severity of the particular disease undergoing therapy.

For administration to non-human animals, the composition can also be added to the animal feed or drinking water. It can be convenient to formulate the animal feed and

5

10

15

20

25

drinking water compositions so that the animal takes in a therapeutically appropriate quantity of the composition along with its diet. It can also be convenient to present the composition as a premix for addition to the feed or drinking water.

The nucleic acid molecules of the present invention can also be administered to a subject in combination with other therapeutic compounds to increase the overall therapeutic effect. The use of multiple compounds to treat an indication can increase the beneficial effects while reducing the presence of side effects.

In one embodiment, the invention comprises compositions suitable for administering nucleic acid molecules of the invention to specific cell types. For example, the asialoglycoprotein receptor (ASGPr) (Wu and Wu, 1987, J. Biol. Chem. 262, 4429-4432) is unique to hepatocytes and binds branched galactose-terminal glycoproteins, such as asialoorosomucoid (ASOR). In another example, the folate receptor is overexpressed in many cancer cells. Binding of such glycoproteins, synthetic glycoconjugates, or folates to the receptor takes place with an affinity that strongly depends on the degree of branching of the oligosaccharide chain, for example, triatennary structures are bound with greater affinity than biatenarry or monoatennary chains (Baenziger and Fiete, 1980, Cell, 22, 611-620; Connolly et al., 1982, J. Biol. Chem., 257, 939-945). Lee and Lee, 1987, Glycoconjugate J., 4, 317-328, obtained this high specificity through the use of N-acetyl-D-galactosamine as the carbohydrate moiety, which has higher affinity for the receptor, compared to galactose. This "clustering effect" has also been described for the binding and uptake of mannosyl-terminating glycoproteins or glycoconjugates (Ponpipom et al., 1981, J. Med. Chem., 24, 1388-1395). The use of galactose, galactosamine, or folate based conjugates to transport exogenous compounds across cell membranes can provide a targeted delivery approach to, for example, the treatment of liver disease, cancers of the liver, or other cancers. The use of bioconjugates can also provide a reduction in the required dose of therapeutic compounds required for treatment. Furthermore, therapeutic bioavialability, pharmacodynamics, and pharmacokinetic parameters can be modulated through the use of nucleic acid bioconjugates of the invention. Non-limiting examples of such bioconjugates are described in Vargeese et al., USSN 10/201,394, filed August 13, 2001; and Matulic-Adamic et al., USSN 60/362,016, filed March 6, 2002.

5

10.

15

20

25

Alternatively, certain siNA molecules of the instant invention can be expressed within cells from eukaryotic promoters (e.g., Izant and Weintraub, 1985, Science, 229, 345; McGarry and Lindquist, 1986, Proc. Natl. Acad. Sci., USA 83, 399; Scanlon et al., 1991, Proc. Natl. Acad. Sci. USA, 88, 10591-5; Kashani-Sabet et al., 1992, Antisense Res. Dev., 2, 3-15; Dropulic et al., 1992, J. Virol., 66, 1432-41; Weerasinghe et al., 1991, J. Virol., 65, 5531-4; Ojwang et al., 1992, Proc. Natl. Acad. Sci. USA, 89, 10802-6; Chen et al., 1992, Nucleic Acids Res., 20, 4581-9; Sarver et al., 1990 Science, 247, 1222-1225; Thompson et al., 1995, Nucleic Acids Res., 23, 2259; Good et al., 1997, Gene Therapy, 4, 45. Those skilled in the art realize that any nucleic acid can be expressed in eukaryotic cells from the appropriate DNA/RNA vector. The activity of such nucleic acids can be augmented by their release from the primary transcript by a enzymatic nucleic acid (Draper et al., PCT WO 93/23569, and Sullivan et al., PCT WO 94/02595; Ohkawa et al., 1992, Nucleic Acids Symp. Ser., 27, 15-6; Taira et al., 1991, Nucleic Acids Res., 19, 5125-30; Ventura et al., 1993, Nucleic Acids Res., 21, 3249-55; Chowrira et al., 1994, J. Biol. Chem., 269, 25856.

In another aspect of the invention, RNA molecules of the present invention can be expressed from transcription units (see for example Couture et al., 1996, TIG., 12, 510) inserted into DNA or RNA vectors. The recombinant vectors can be DNA plasmids or viral vectors siNA expressing viral vectors can be constructed based on, but not limited to, adeno-associated virus, retrovirus, adenovirus, or alphavirus. In another embodiment, pol III based constructs are used to express nucleic acid molecules of the invention (see for example Thompson, U.S. Pats. Nos. 5,902,880 and 6,146,886). The recombinant vectors capable of expressing the siNA molecules can be delivered as described above, and persist in target cells. Alternatively, viral vectors can be used that provide for transient expression of nucleic acid molecules. Such vectors can be repeatedly administered as necessary. Once expressed, the siNA molecule interacts with the target mRNA and generates an RNAi response. Delivery of siNA molecule expressing vectors can be systemic, such as by intravenous or intra-muscular administration, by administration to target cells ex-planted from a subject followed by reintroduction into the subject, or by any other means that would allow for introduction into the desired target cell (for a review see Couture et al., 1996, TIG., 12, 510).

5

10

15

20

25

In one aspect the invention features an expression vector comprising a nucleic acid sequence encoding at least one siNA molecule of the instant invention. The expression vector can encode one or both strands of a siNA duplex, or a single self-complementary strand that self hybridizes into a siNA duplex. The nucleic acid sequences encoding the siNA molecules of the instant invention can be operably linked in a manner that allows expression of the siNA molecule (see for example Paul et al., 2002, Nature Biotechnology, 19, 505; Miyagishi and Taira, 2002, Nature Biotechnology, 19, 497; Lee et al., 2002, Nature Biotechnology, 19, 500; and Novina et al., 2002, Nature Medicine, advance online publication doi:10.1038/nm725).

In another aspect, the invention features an expression vector comprising: a) a transcription initiation region (e.g., eukaryotic pol I, II or III initiation region); b) a transcription termination region (e.g., eukaryotic pol I, II or III termination region); and c) a nucleic acid sequence encoding at least one of the siNA molecules of the instant invention; wherein said sequence is operably linked to said initiation region and said termination region, in a manner that allows expression and/or delivery of the siNA molecule. The vector can optionally include an open reading frame (ORF) for a protein operably linked on the 5' side or the 3'-side of the sequence encoding the siNA of the invention; and/or an intron (intervening sequences).

Transcription of the siNA molecule sequences can be driven from a promoter for eukaryotic RNA polymerase I (pol I), RNA polymerase II (pol II), or RNA polymerase III (pol III). Transcripts from pol II or pol III promoters are expressed at high levels in all cells; the levels of a given pol II promoter in a given cell type depends on the nature of the gene regulatory sequences (enhancers, silencers, etc.) present nearby. Prokaryotic RNA polymerase promoters are also used, providing that the prokaryotic RNA polymerase enzyme is expressed in the appropriate cells (Elroy-Stein and Moss, 1990, *Proc. Natl. Acad. Sci. U S A*, 87, 6743-7; Gao and Huang 1993, *Nucleic Acids Res.*, 21, 2867-72; Lieber *et al.*, 1993, *Methods Enzymol.*, 217, 47-66; Zhou *et al.*, 1990, *Mol. Cell. Biol.*, 10, 4529-37). Several investigators have demonstrated that nucleic acid molecules expressed from such promoters can function in mammalian cells (e.g. Kashani-Sabet *et al.*, 1992, *Antisense Res. Dev.*, 2, 3-15; Ojwang *et al.*, 1992, *Proc. Natl. Acad. Sci. U S A*, 89, 10802-6; Chen *et al.*, 1992, *Nucleic Acids Res.*, 20, 4581-9; Yu *et al.*, 1993, *Proc. Natl. Acad. Sci. U S A*, 90, 6340-4; L'Huillier *et al.*, 1992, *EMBO J.*, 11,

5

10

15

20

25

4411-8; Lisziewicz et al., 1993, Proc. Natl. Acad. Sci. U. S. A, 90, 8000-4; Thompson et al., 1995, Nucleic Acids Res., 23, 2259; Sullenger & Cech, 1993, Science, 262, 1566). More specifically, transcription units such as the ones derived from genes encoding U6 small nuclear (snRNA), transfer RNA (tRNA) and adenovirus VA RNA are useful in generating high concentrations of desired RNA molecules such as siNA in cells (Thompson et al., supra; Couture and Stinchcomb, 1996, supra; Noonberg et al., 1994, Nucleic Acid Res., 22, 2830; Noonberg et al., U.S. Pat. No. 5,624,803; Good et al., 1997, Gene Ther., 4, 45; Beigelman et al., International PCT Publication No. WO 96/18736. The above siNA transcription units can be incorporated into a variety of vectors for introduction into mammalian cells, including but not restricted to, plasmid DNA vectors, viral DNA vectors (such as adenovirus or adeno-associated virus vectors), or viral RNA vectors (such as retroviral or alphavirus vectors) (for a review see Couture and Stinchcomb, 1996, supra).

In another aspect the invention features an expression vector comprising a nucleic acid sequence encoding at least one of the siNA molecules of the invention in a manner that allows expression of that siNA molecule. The expression vector comprises in one embodiment; a) a transcription initiation region; b) a transcription termination region; and c) a nucleic acid sequence encoding at least one strand of the siNA molecule, wherein the sequence is operably linked to the initiation region and the termination region in a manner that allows expression and/or delivery of the siNA molecule.

In another embodiment the expression vector comprises: a) a transcription initiation region; b) a transcription termination region; c) an open reading frame; and d) a nucleic acid sequence encoding at least one strand of a siNA molecule, wherein the sequence is operably linked to the 3'-end of the open reading frame and wherein the sequence is operably linked to the initiation region, the open reading frame and the termination region in a manner that allows expression and/or delivery of the siNA molecule. In yet another embodiment, the expression vector comprises: a) a transcription initiation region; b) a transcription termination region; c) an intron; and d) a nucleic acid sequence encoding at least one siNA molecule, wherein the sequence is operably linked to the initiation region, the intron and the termination region in a manner which allows expression and/or delivery of the nucleic acid molecule.

5

10

15

20

25

In another embodiment, the expression vector comprises: a) a transcription initiation region; b) a transcription termination region; c) an intron; d) an open reading frame; and e) a nucleic acid sequence encoding at least one strand of a siNA molecule, wherein the sequence is operably linked to the 3'-end of the open reading frame and wherein the sequence is operably linked to the initiation region, the intron, the open reading frame and the termination region in a manner which allows expression and/or delivery of the siNA molecule.

### Examples:

5

15

20

25

The following are non-limiting examples showing the selection, isolation, synthesis and activity of nucleic acids of the instant invention.

## Example 1: Tandem synthesis of siNA constructs

Exemplary siNA molecules of the invention are synthesized in tandem using a cleavable linker, for example, a succinyl-based linker. Tandem synthesis as described herein is followed by a one-step purification process that provides RNAi molecules in high yield. This approach is highly amenable to siNA synthesis in support of high throughput RNAi screening, and can be readily adapted to multi-column or multi-well synthesis platforms.

After completing a tandem synthesis of a siNA oligo and its complement in which the 5'-terminal dimethoxytrityl (5'-O-DMT) group remains intact (trityl on synthesis), the oligonucleotides are deprotected as described above. Following deprotection, the siNA sequence strands are allowed to spontaneously hybridize. This hybridization yields a duplex in which one strand has retained the 5'-O-DMT group while the complementary strand comprises a terminal 5'-hydroxyl. The newly formed duplex behaves as a single molecule during routine solid-phase extraction purification (Trityl-On purification) even though only one molecule has a dimethoxytrityl group. Because the strands form a stable duplex, this dimethoxytrityl group (or an equivalent group, such as other trityl groups or other hydrophobic moieties) is all that is required to purify the pair of oligos, for example, by using a C18 cartridge.

Standard phosphoramidite synthesis chemistry is used up to the point of introducing a tandem linker, such as an inverted deoxy abasic succinate or glyceryl succinate linker (see Figure 1) or an equivalent cleavable linker. A non-limiting example of linker coupling conditions that can be used includes a hindered base such as diisopropylethylamine (DIPA) and/or DMAP in the presence of an activator reagent such as Bromotripyrrolidinophosphoniumhexaflurorophosphate (PyBrOP). After the linker is coupled, standard synthesis chemistry is utilized to complete synthesis of the second sequence leaving the terminal the 5'-O-DMT intact. Following synthesis, the resulting oligonucleotide is deprotected according to the procedures described herein and quenched with a suitable buffer, for example with 50mM NaOAc or 1.5M NH4H2CO3.

Purification of the siNA duplex can be readily accomplished using solid phase extraction, for example using a Waters C18 SepPak 1g cartridge conditioned with 1 column volume (CV) of acetonitrile, 2 CV H2O, and 2 CV 50mM NaOAc. The sample is loaded and then washed with 1 CV H2O or 50mM NaOAc. Failure sequences are eluted with 1 CV 14% ACN (Aqueous with 50mM NaOAc and 50mM NaCl). The column is then washed, for example with 1 CV H2O followed by on-column detritylation, for example by passing 1 CV of 1% aqueous trifluoroacetic acid (TFA) over the column, then adding a second CV of 1% aqueous TFA to the column and allowing to stand for approximately 10 minutes. The remaining TFA solution is removed and the column washed with H2O followed by 1 CV 1M NaCl and additional H2O. The siNA duplex product is then eluted, for example, using 1 CV 20% aqueous CAN.

Figure 2 provides an example of MALDI-TOV mass spectrometry analysis of a purified siNA construct in which each peak corresponds to the calculated mass of an individual siNA strand of the siNA duplex. The same purified siNA provides three peaks when analyzed by capillary gel electrophoresis (CGE), one peak presumably corresponding to the duplex siNA, and two peaks presumably corresponding to the separate siNA sequence strands. Ion exchange HPLC analysis of the same siNA contract only shows a single peak. Testing of the purified siNA construct using a luciferase reporter assay described below demonstrated the same RNAi activity compared to siNA constructs generated from separately synthesized oligonucleotide sequence strands.

Example 2: Serum stability of chemically modified siNA constructs

5

10

15

20

25

Chemical modifications were introduced into siNA constructs to determine the stability of these constructs compared to native siNA oligonucleotides (containing two thymidine nucleotide overhangs) in human serum. An investigation of the serum stability of RNA duplexes revealed that siNA constructs consisting of all RNA nucleotides containing two thymidine nucleotide overhangs have a half-life in serum of 15 seconds, whereas chemically modified siNA constructs remained stable in serum for 1 to 3 days depending on the extent of modification. RNAi stability tests were performed by internally labeling one strand (strand 1) of siNA and duplexing with 1.5 X the concentration of the complementary siNA strand (strand 2) (to insure all labeled material was in duplex form). Duplexed siNA constructs were then tested for stability by incubating at a final concentration of 2µM siNA (strand 2 concentration) in 90% mouse or human serum for time-points of 30sec, 1min, 5min, 30min, 90min, 4hrs 10min, 16hrs 24min, and 49hrs. Time points were run on a 15% denaturing polyacrylamide gels and analyzed on a phosphoimager.

5

10

15

20

25

30

BNSDOCID: <WO

\_03074654A2\_I\_>

Internal labeling was performed via kinase reactions with polynucleotide kinase (PNK) and <sup>32</sup>P-γ-ATP, with addition of radiolabeled phosphate at nucleotide 13 of strand 2, counting in from the 3' side. Ligation of the remaining 8-mer fragments with T4 RNA ligase resulted in the full length, 21-mer, strand 2. Duplexing of RNAi was done by adding appropriate concentrations of the siNA oligonucleotides and heating to 95° C for 5min followed by slow cooling to room temperature. Reactions were performed by adding 100% serum to the siNA duplexes and incubating at 37° C, then removing aliquots at desired time-points. Results of this study are summarized in **Figure 3**. As shown in the Figure 3, chemically modified siNA molecules (e.g., SEQ ID NOs: 925/927, 925/928, 925/929, 925/930, and 925/931) have significantly increased serum stability compared to an siNA construct having all ribonucleotides except a 3'-terminal dithymidine (TT) modification (e.g., SEQ ID NOs: 925/926).

# Example 3: Identification of potential siNA target sites in any RNA sequence

The sequence of an RNA target of interest, such as a viral or human mRNA transcript, is screened for target sites, for example by using a computer folding algorithm. In a non-limiting example, the sequence of a gene or RNA gene transcript derived from a database, such as Genbank, is used to generate siNA targets having complementarity to

the target. Such sequences can be obtained from a database, or can be determined experimentally as known in the art. Target sites that are known, for example, those target sites determined to be effective target sites based on studies with other nucleic acid molecules, for example ribozymes or antisense, or those targets known to be associated with a disease or condition such as those sites containing mutations or deletions, can be used to design siNA molecules targeting those sites. Various parameters can be used to determine which sites are the most suitable target sites within the target RNA sequence. These parameters include but are not limited to secondary or tertiary RNA structure, the nucleotide base composition of the target sequence, the degree of homology between various regions of the target sequence, or the relative position of the target sequence within the RNA transcript. Based on these determinations, any number of target sites within the RNA transcript can be chosen to screen siNA molecules for efficacy, for example by using in vitro RNA cleavage assays, cell culture, or animal models. In a nonlimiting example, anywhere from 1 to 1000 target sites are chosen within the transcript based on the size of the siNA construct to be used. High throughput screening assays can be developed for screening siNA molecules using methods known in the art, such as with multi-well or multi-plate assays or combinatorial/siNA library screening assays to determine efficient reduction in target gene expression.

#### Example 4: Selection of siNA molecule target sites in a RNA

The following non-limiting steps can be used to carry out the selection of siNAs targeting a given gene sequence or transcript.

The target sequence is parsed in silico into a list of all fragments or subsequences of a particular length, for example 23 nucleotide fragments, contained within the target sequence. This step is typically carried out using a custom Perl script, but commercial sequence analysis programs such as Oligo, MacVector, or the GCG Wisconsin Package can be employed as well.

In some instances the siNAs correspond to more than one target sequence; such would be the case for example in targeting different transcripts of the same gene, targeting different transcripts of more than one gene, or for targeting both the human gene and an animal homolog. In this case, a subsequence list of a particular length is generated for each of the targets, and then the lists are compared to find matching sequences in each

5

10

15

20

25

list. The subsequences are then ranked according to the number of target sequences that contain the given subsequence; the goal is to find subsequences that are present in most or all of the target sequences. Alternately, the ranking can identify subsequences that are unique to a target sequence, such as a mutant target sequence. Such an approach would enable the use of siNA to target specifically the mutant sequence and not effect the expression of the normal sequence.

In some instances the siNA subsequences are absent in one or more sequences while present in the desired target sequence; such would be the case if the siNA targets a gene with a paralogous family member that is to remain untargeted. As in case 2 above, a subsequence list of a particular length is generated for each of the targets, and then the lists are compared to find sequences that are present in the target gene but are absent in the untargeted paralog.

The ranked siNA subsequences can be further analyzed and ranked according to GC content. A preference can be given to sites containing 30-70% GC, with a further preference to sites containing 40-60% GC.

The ranked siNA subsequences can be further analyzed and ranked according to self-folding and internal hairpins. Weaker internal folds are preferred; strong hairpin structures are to be avoided.

The ranked siNA subsequences can be further analyzed and ranked according to whether they have runs of GGG or CCC in the sequence. GGG (or even more Gs) in either strand can make oligonucleotide synthesis problematic and can potentially interfere with RNAi activity, so it is avoided whenever other appropriately suitable sequences are available. CCC is searched in the target strand because that will place GGG in the antisense strand.

The ranked siNA subsequences can be further analyzed and ranked according to whether they have the dinucleotide UU (uridine dinucleotide) on the 3'-end of the sequence, and/or AA on the 5'-end of the sequence (to yield 3' UU on the antisense sequence). These sequences allow one to design siNA molecules with terminal TT thymidine dinucleotides.

5

10

15

20

Four or five target sites are chosen from the ranked list of subsequences as described above. For example, in subsequences having 23 nucleotides, the right 21 nucleotides of each chosen 23-mer subsequence are then designed and synthesized for the upper (sense) strand of the siNA duplex, while the reverse complement of the left 21 nucleotides of each chosen 23-mer subsequence are then designed and synthesized for the lower (antisense) strand of the siNA duplex (see Tables I). If terminal TT residues are desired for the sequence (as described in paragraph 7), then the two 3' terminal nucleotides of both the sense and antisense strands are replaced by TT prior to synthesizing the oligos.

The siNA molecules are screened in an in vitro, cell culture or animal model system to identify the most active siNA molecule or the most preferred target site within the target RNA sequence.

In an alternate approach, a pool of siNA constructs specific to a target sequence is used to screen for target sites in cells expressing target RNA, such as human HeLa cells. The general strategy used in this approach is shown in Figure 21. A non-limiting example of such as pool is a pool comprising sequences having antisense sequences complementary to the target RNA sequence and sense sequences complementary to the antisense sequences. Cells (e.g., HeLa cells) expressing the target gene are transfected with the pool of siNA constructs and cells that demonstrate a phenotype associated with gene silencing are sorted. The pool of siNA constructs can be chemically modified as described herein and synthesized, for example, in a high throughput manner. The siNA from cells demonstrating a positive phenotypic change (e.g., decreased target mRNA levels or target protein expression), are identified, for example by positional analysis within the assay, and are used to determine the most suitable target site(s) within the target RNA sequence based upon the complementary sequence to the corresponding siNA antisense strand identified in the assay.

## Example 5: RNAi activity of chemically modified siNA constructs

Short interfering nucleic acid (siNA) is emerging as a powerful tool for gene regulation. All-ribose siNA duplexes activate the RNAi pathway but have limited utility as therapeutic compounds due to their nuclease sensitivity and short half-life in serum, as shown in Example 2 above. To develop nuclease-resistant siNA constructs for *in vivo* 

5

10

15

20

25

applications, siNAs that target luciferase mRNA and contain stabilizing chemical modifications were tested for activity in HeLa cells. The sequences for the siNA oligonucleotide sequences used in this study are shown in **Table I**. Modifications included phosphorothioate linkages (P=S), 2'-O-methyl nucleotides, or 2'-fluoro (F) nucleotides in one or both siNA strands and various 3'-end stabilization chemistries, including 3'-glyceryl, 3'-inverted abasic, 3'-inverted Thymidine, and/or Thymidine. Active siNA containing stabilizing modifications such as described herein should prove useful for *in vivo* applications.

A luciferase reporter system was utilized to test RNAi activity of chemically modified siNA constructs compared to siNA constructs consisting of all RNA nucleotides containing two thymidine nucleotide overhangs. Sense and antisense siNA strands (20 uM each) were annealed by incubation in buffer (100 mM potassium acetate, 30 mM HEPES-KOH, pH 7.4, 2 mM magnesium acetate) for 1 min. at 90°C followed by 1 hour at 37°C. Plasmids encoding firefly luciferase (pGL2) and renilla luciferase (pRLSV40) were purchased from Promega Biotech.

HeLa S3 cells were grown at 37°C in DMEM with 5% FBS and seeded at 15,300 cells in 100 ul media per well of a 96-well plate 24 hours prior to transfection. For transfection, 4 ul Lipofectamine 2000 (Life Technologies) was added to 96 ul OPTI-MEM, vortexed and incubated at room temperature for 5 minutes. The 100 ul diluted lipid was then added to a microtiter tube containing 5 ul pGL2 (200ng/ul), 5 ul pRLSV40 (8 ng/ul) 6 ul siNA (25 nM or 10 nM final), and 84 ul OPTI-MEM, vortexed briefly and incubated at room temperature for 20 minutes. The transfection mix was then mixed briefly and 50 ul was added to each of three wells that contained HeLa S3 cells in 100 ul media. Cells were incubated for 20 hours after transfection and analyzed for luciferase expression using the Dual luciferase assay according to the manufacturer's instructions (Promega Biotech). The results of this study are summarized in Figures 4-16. sequences of the siNA strands used in this study are shown in Table I and are referred to by RPI# in the figures. Normalized luciferase activity is reported as the ratio of firefly luciferase activity to renilla luciferase activity in the same sample. Error bars represent standard deviation of triplicate transfections. As shown in Figures 4-16, the RNAi activity of chemically modified constructs is comparable to that of control siNA constructs, which consist of all ribonucleotides at every position except the 3'-terminus

5

10

15

20

25

which comprises two thymidine nucleotide overhangs. In some instances, the RNAi activity of the chemically modified constructs is greater than the siNA construct consisting of all ribonucleotides at every position except the 3'-terminus which comprises two thymidine nucleotide overhangs. For example, **Figure 4** shows results obtained from a screen using phosphorothioate modified siNA constructs; the RPI 27654/27659 construct contains phosphorothioate substitutions for every pyrimidine nucleotide in both sequences, the RPI 27657/27662 construct contains 5 terminal 3'-phosphorothioate substitutions in each strand, the RPI 27649/27658 construct contains all phosphorothioate substitutions only in the antisense strand, whereas the RPI 27649/27660 and RPI 27649/27661 constructs have unmodified sense strands and varying degrees of phosphorothioate substitutions in the antisense strand. All of these constructs show significant RNAi activity when compared to a scrambled siNA.

Figure 5 shows results obtained from a screen using phosphorothioate (RPI 28253/28255 and RPI 28254/28256) and universal base substitutions (RPI 28257/28259 and RPI 28258/28260) compared to the same controls described above. As shown, these modifications show equivalent or better RNAi activity when compared to the control siNA construct.

**Figure 6** shows results obtained from a screen using 2'-O-methyl modified siNA constructs in which the sense strand contains either 10 (RPI 28244/27650) or 5 (RPI 28245/27650) 2'-O-methyl substitutions, both with comparable activity to the control siNA construct.

Figure 7 shows results obtained from a screen using 2'-O-methyl or 2'-deoxy-2'-fluoro modified siNA constructs compared to a control construct consisting of all ribonucleotides at every position except the 3'-terminus which comprises two thymidine nucleotide overhangs.

Figure 8 compares a siNA construct containing six phosphorothioate substitutions in each strand (RPI 28460/28461), where 5 phosphorothioates are present at the 3' end and a single phosphorothioate is present at the 5' end of each strand. This motif shows very similar activity to the control siNA construct consisting of all ribonucleotides at every position except the 3'-terminus which comprises two thymidine nucleotide overhangs.

5

10

15

20

25

Figure 9 compares a siNA construct synthesized by the method of the invention described in Example 1, wherein an inverted deoxyabasic succinate linker was used to generate a siNA having a 3'-inverted deoxyabasic cap on the antisense strand of the siNA. This construct shows improved activity compared to the control siNA (siGL2) construct consisting of all ribonucleotides at every position except the 3'-terminus which comprises two thymidine nucleotide overhangs.

Figure 10 shows the results of an RNAi activity screen of chemically modified siNA constructs including 3'-glyceryl modified siNA constructs compared to an all RNA control siNA construct using a luciferase reporter system. These chemically modified siNAs were compared in the luciferase assay described herein at 1 nM and 10nM concentration using an all RNA siNA control (siGL2) having having 3'-terminal dithymidine (TT) and its corresponding inverted control (Inv siGL2). The background level of luciferase expression in the HeLa cells is designated by the "cells" column. Sense and antisense strands of chemically modified siNA constructs are shown by RPI number (sense strand/antisense strand). Sequences correspoding to these RPI numbers are shown in Table I. As shown in the Figure, the 3'-terminal modified siNA constructs retain significant RNAi activity compared to the control siNA (siGL2) construct.

Figure 11 shows the results of an RNAi activity screen of chemically modified siNA constructs. The screen compared various combinations of sense strand chemical modifications and antisense strand chemical modifications. These chemically modified siNAs were compared in the luciferase assay described herein at 1 nM and 10nM concentration using an all RNA siNA control (siGL2) having having 3'-terminal dithymidine (TT) and its corresponding inverted control (Inv siGL2). The background level of luciferase expression in the HeLa cells is designated by the "cells" column. Sense and antisense strands of chemically modified siNA constructs are shown by RPI number (sense strand/antisense strand). Sequences correspoding to these RPI numbers are shown in Table I. As shown in the figure, the chemically modified RPI 30063/30430, RPI 30433/30430, and RPI 30063/30224 constructs retain significant RNAi activity compared to the control siNA construct. It should be noted that RPI 30433/30430 is a siNA construct having no ribonucleotides which retains significant RNAi activity compared to the constrol siGL2 construct in vitro, therefore, this construct is expected to

5

10

15

20

25

have both similar RNAi activity and improved stability compared to siNA constructs having ribonucleotides in vivo.

Figure 12 shows the results of an RNAi activity screen of chemically modifed siNA constructs. The screen compared various combinations of sense strand chemical modifications and antisense strand chemical modifications. These chemically modified siNAs were compared in the luciferase assay described herein at 1 nM and 10nM concentration using an all RNA siNA control (siGL2) having having 3'-terminal dithymidine (TT) and its corresponding inverted control (Inv siGL2). The background level of luciferase expression in the HeLa cells is designated by the "cells" column. Sense and antisense strands of chemically modified siNA constructs are shown by RPI number (sense strand/antisense strand). Sequences correspoding to these RPI numbers are shown in Table I. As shown in the figure, the chemically modified RPI 30063/30224 and RPI 30063/30430 constructs retain significant RNAi activity compared to the control siNA (siGL2) construct. In addition, the antisense strand alone (RPI 30430) and an inverted control (RPI 30227/30229, having matched chemistry to RPI 30063/30224) were compared to the siNA duplexes described above. The antisense strand (RPI 30430) alone provides far less inhibition compared to the siNA duplexes using this sequence.

Figure 13 shows the results of an RNAi activity screen of chemically modified siNA constructs. The screen compared various combinations of sense strand chemical modifications and antisense strand chemical modifications. These chemically modified siNAs were compared in the luciferase assay described herein at 1 nM and 10nM concentration using an all RNA siNA control (siGL2) having having 3'-terminal dithymidine (TT) and its corresponding inverted control (Inv siGL2). The background level of luciferase expression in the HeLa cells is designated by the "cells" column. Sense and antisense strands of chemically modified siNA constructs are shown by RPI number (sense strand/antisense strand). Sequences correspoding to these RPI numbers are shown in Table I. In addition, an inverted control (RPI 30226/30229, having matched chemistry to RPI 30222/30224) was compared to the siNA duplexes described above. As shown in the figure, the chemically modified RPI 28251/30430, RPI 28251/30224, and RPI 30222/30224 constructs retain significant RNAi activity compared to the control siNA construct, and the chemically modified RPI 28251/30430 construct demonstrates improved activity compared to the control siNA (siGL2) construct.

5

10

15

20

25

Figure 14 shows the results of an RNAi activity screen of chemically modified siNA constructs including various 3'-terminal modified siNA constructs compared to an all RNA control siNA construct using a luciferase reporter system. These chemically modified siNAs were compared in the luciferase assay described herein at 1 nM and 10nM concentration using an all RNA siNA control (siGL2) having having 3'-terminal dithymidine (TT) and its corresponding inverted control (Inv siGL2). The background level of luciferase expression in the HeLa cells is designated by the "cells" column. Sense and antisense strands of chemically modified siNA constructs are shown by RPI number (sense strand/antisense strand). Sequences correspoding to these RPI numbers are shown in Table I. As shown in the figure, the chemically modified RPI 30222/30546, 30222/30224, 30222/30551, 30222/30557 and 30222/30558 constructs retain significant RNAi activity compared to the control siNA construct.

Figure 15 shows the results of an RNAi activity screen of chemically modified siNA constructs. The screen compared various combinations of sense strand chemistries compared to a fixed antisense strand chemistry. These chemically modified siNAs were compared in the luciferase assay described herein at 1 nM and 10nM concentration using an all RNA siNA control (siGL2) having having 3'-terminal dithymidine (TT) and its corresponding inverted control (Inv siGL2). The background level of luciferase expression in the HeLa cells is designated by the "cells" column. Sense and antisense strands of chemically modified siNA constructs are shown by RPI number (sense strand/antisense strand). Sequences correspoding to these RPI numbers are shown in Table I. As shown in the figure, the chemically modified RPI 30063/30430, 30434/30430, and 30435/30430 constructs all demonstrate greater activity compared to the control siNA (siGL2) construct.

## Example 6: RNAi activity titration

A titration assay was performed to determine the lower range of siNA concentration required for RNAi activity both in a control siNA construct consisting of all RNA nucleotides containing two thymidine nucleotide overhangs and a chemically modified siNA construct comprising 5 phosphorothioate internucleotide linkages in both the sense and antisense strands. The assay was performed as described above, however, the siNA constructs were diluted to final concentrations between 2.5 nM and 0.025 nM. Results

5

10

20

25

are shown in Figure 16. As shown in Figure 16, the chemically modified siNA construct shows a very similar concentration dependent RNAi activity profile to the control siNA construct when compared to an inverted siNA sequence control.

### Example 7: siNA design

5

10

15

20

25

30

siNA target sites were chosen by analyzing sequences of the target RNA and optionally prioritizing the target sites on the basis of folding (structure of any given sequence analyzed to determine siNA accessibility to the target), by using a library of siNA molecules as described in Example 4, or alternately by using an *in vitro* siNA system as described in Example 9 herein. siNA molecules were designed that could bind each target and are optionally individually analyzed by computer folding to assess whether the siNA molecule can interact with the target sequence. Varying the length of the siNA molecules can be chosen to optimize activity. Generally, a sufficient number of complementary nucleotide bases are chosen to bind to, or otherwise interact with, the target RNA, but the degree of complementarity can be modulated to accommodate siNA duplexes or varying length or base composition. By using such methodologies, siNA molecules can be designed to target sites within any known RNA sequence, for example those RNA sequences corresponding to the any gene transcript.

Chemically modified siNA constructs are designed to provide nuclease stability for systemic administration in vivo and/or improved pharmacokinetic, localization, and delivery properties while preserving the ability to mediate RNAi activity. Chemical modifications as described herein are introduced synthetically using synthetic methods described herein and those generally known in the art. The synthetic siNA constructs are then assayed for nuclease stability in serum and/or cellular/tissue extracts (e.g. liver extracts). The synthetic siNA constructs are also tested in parallel for RNAi activity using an appropriate assay, such as a luciferase reporter assay as described herein or another suitable assay that can quantity RNAi activity. Synthetic siNA constructs that possess both nuclease stability and RNAi activity can be further modified and reevaluated in stability and activity assays. The chemical modifications of the stabilized active siNA constructs can then be applied to any siNA sequence targeting any chosen RNA and used, for example, in target screening assays to pick lead siNA compounds for therapeutic development (see for example Figure 24).

## Example 8: Chemical Synthesis and Purification of siNA

siNA molecules can be designed to interact with various sites in the RNA message, for example, target sequences within the RNA sequences described herein. The sequence of one strand of the siNA molecule(s) is complementary to the target site sequences described above. The siNA molecules can be chemically synthesized using methods described herein. Inactive siNA molecules that are used as control sequences can be synthesized by scrambling the sequence of the siNA molecules such that it is not complementary to the target sequence. Generally, siNA constructs can by synthesized using solid phase oligonucleotide synthesis methods as described herein (see for example Usman *et al.*, US Patent Nos. 5,804,683; 5,831,071; 5,998,203; 6,117,657; 6,353,098; 6,362,323; 6,437,117; 6,469,158; Scaringe *et al.*, US Patent Nos. 6,111,086; 6,008,400; 6,111,086 all incorporated by reference herein in their entirety).

In a non-limiting example, RNA oligonucleotides are synthesized in a stepwise fashion using the phosphoramidite chemistry as is known in the art. Standard phosphoramidite chemistry involves the use of nucleosides comprising any of 5'-O-dimethoxytrityl, 2'-O-tert-butyldimethylsilyl, 3'-O-2-Cyanoethyl N,N-diisopropylphosphoroamidite groups, and exocyclic amine protecting groups (e.g. N6-benzoyl adenosine, N4 acetyl cytidine, and N2-isobutyryl guanosine). Alternately, 2'-O-Silyl Ethers can be used in conjunction with acid-labile 2'-O-orthoester protecting groups in the synthesis of RNA as described by Scaringe *supra*. Differing 2' chemistries can require different protecting groups, for example 2'-deoxy-2'-amino nucleosides can utilize N-phthaloyl protection as described by Usman *et al.*, US Patent 5,631,360, incorporated by reference herein in its entirety).

During solid phase synthesis, each nucleotide is added sequentially (3'- to 5'-direction) to the solid support-bound oligonucleotide. The first nucleoside at the 3'-end of the chain is covalently attached to a solid support (e.g., controlled pore glass or polystyrene) using various linkers. The nucleotide precursor, a ribonucleoside phosphoramidite, and activator are combined resulting in the coupling of the second nucleoside phosphoramidite onto the 5'-end of the first nucleoside. The support is then washed and any unreacted 5'-hydroxyl groups are capped with a capping reagent such as acetic anhydride to yield inactive 5'-acetyl moieties. The trivalent phosphorus linkage is

5

10

15

20

25

then oxidized to a more stable phosphate linkage. At the end of the nucleotide addition cycle, the 5'-O-protecting group is cleaved under suitable conditions (e.g., acidic conditions for trityl-based groups and Fluoride for silyl-based groups). The cycle is repeated for each subsequent nucleotide.

Modification of synthesis conditions can be used to optimize coupling efficiency, for example by using differing coupling times, differing reagent/phosphoramidite concentrations, differing contact times, differing solid supports and solid support linker chemistries depending on the particular chemical composition of the siNA to be synthesized. Deprotection and purification of the siNA can be performed as is generally described in Usman et al., US 5,831,071, US 6,353,098, US 6,437,117, and Bellon et al., US 6,054,576, US 6,162,909, US 6,303,773, incorporated by reference herein in their entirety or Scaringe *supra*,. Additionally, deprotection conditions can be modified to provide the best possible yield and purity of siNA constructs. For example, applicant has observed that oligonucleotides comprising 2'-deoxy-2'-fluoro nucleotides can degrade under inappropriate deprotection conditions. Such oligonucleotides are deprotected using aqueous methylamine at about 35°C for 30 minutes. If the 2'-deoxy-2'-fluoro containing oligonucleotide also comprises ribonucleotides, after deprotection with aqueous methylamine at about 35°C for 30 minutes, TEA-HF is added and the reaction maintained at about 65°C for an additional 15 minutes.

### 20 Example 9: RNAi in vitro assay to assess siNA activity

An in vitro assay that recapitulates RNAi in a cell free system is used to evaluate siNA constructs specific to target RNA. The assay comprises the system described by Tuschl et al., 1999, Genes and Development, 13, 3191-3197 and Zamore et al., 2000, Cell, 101, 25-33 adapted for use with target RNA. A Drosophila extract derived from syncytial blastoderm is used to reconstitute RNAi activity in vitro. Target RNA is generated via in vitro transcription from an appropriate plasmid using T7 RNA polymerase or via chemical synthesis as described herein. Sense and antisense siNA strands (for example 20 uM each) are annealed by incubation in buffer (such as 100 mM potassium acetate, 30 mM HEPES-KOH, pH 7.4, 2 mM magnesium acetate) for 1 min. at 90°C followed by 1 hour at 37°C, then diluted in lysis buffer (for example 100 mM potassium acetate, 30 mM HEPES-KOH at pH 7.4, 2mM magnesium acetate). Annealing

5

10

15

25

can be monitored by gel electrophoresis on an agarose gel in TBE buffer and stained with ethidium bromide. The Drosophila lysate is prepared using zero to two-hour-old embryos from Oregon R flies collected on yeasted molasses agar that are dechorionated and lysed. The lysate is centrifuged and the supernatant isolated. The assay comprises a reaction mixture containing 50% lysate [vol/vol], RNA (10-50 pM final concentration), and 10% [vol/vol] lysis buffer containing siNA (10 nM final concentration). The reaction mixture also contains 10 mM creatine phosphate, 10 ug.ml creatine phosphokinase, 100 um GTP, 100 uM UTP, 100 uM CTP, 500 uM ATP, 5 mM DTT, 0.1 U/uL RNasin (Promega), and 100 uM of each amino acid. The final concentration of potassium acetate is adjusted to 100 mM. The reactions are pre-assembled on ice and preincubated at 25° C for 10 minutes before adding RNA, then incubated at 25° C for an additional 60 minutes. Reactions are quenched with 4 volumes of 1.25 x Passive Lysis Buffer (Promega). Target RNA cleavage is assayed by RT-PCR analysis or other methods known in the art and are compared to control reactions in which siNA is omitted from the reaction.

Alternately, internally-labeled target RNA for the assay is prepared by *in vitro* transcription in the presence of [alpha-<sup>32</sup>p] CTP, passed over a G 50 Sephadex column by spin chromatography and used as target RNA without further purification. Optionally, target RNA is 5'-<sup>32</sup>P-end labeled using T4 polynucleotide kinase enzyme. Assays are performed as described above and target RNA and the specific RNA cleavage products generated by RNAi are visualized on an autoradiograph of a gel. The percentage of cleavage is determined by Phosphor Imager<sup>®</sup> quantitation of bands representing intact control RNA or RNA from control reactions without siNA and the cleavage products generated by the assay.

In one embodiment, this assay is used to determine target sites the RNA target for siNA mediated RNAi cleavage, wherein a plurality of siNA constructs are screened for RNAi mediated cleavage of the RNA target, for example, by analyzing the assay reaction by electrophoresis of labeled target RNA, or by northern blotting, as well as by other methodology well known in the art.

## Example 10: Nucleic acid inhibition of target RNA in vivo

5

10

15

20

siNA molecules targeted to the target RNA are designed and synthesized as described above. These nucleic acid molecules can be tested for cleavage activity *in vivo*, for example, using the following procedure.

Two formats are used to test the efficacy of siNAs targeting a particular gene transcipt. First, the reagents are tested on target expressing cells (e.g., HeLa), to determine the extent of RNA and protein inhibition. siNA reagents are selected against the RNA target. RNA inhibition is measured after delivery of these reagents by a suitable transfection agent to cells. Relative amounts of target RNA are measured versus actin using real-time PCR monitoring of amplification (eg., ABI 7700 Taqman®). A comparison is made to a mixture of oligonucleotide sequences made to unrelated targets or to a randomized siNA control with the same overall length and chemistry, but randomly substituted at each position. Primary and secondary lead reagents are chosen for the target and optimization performed. After an optimal transfection agent concentration is chosen, a RNA time-course of inhibition is performed with the lead siNA molecule. In addition, a cell-plating format can be used to determine RNA inhibition.

#### Delivery of siNA to Cells

5

10

15

20

25

30

Cells (e.g., HeLa) are seeded, for example, at 1x10<sup>5</sup> cells per well of a six-well dish in EGM-2 (BioWhittaker) the day before transfection. siNA (final concentration, for example 20nM) and cationic lipid (e.g., final concentration 2µg/ml) are complexed in EGM basal media (Biowhittaker) at 37°C for 30 mins in polystyrene tubes. Following vortexing, the complexed siNA is added to each well and incubated for the times indicated. For initial optimization experiments, cells are seeded, for example, at 1x10<sup>3</sup> in 96 well plates and siNA complex added as described. Efficiency of delivery of siNA to cells is determined using a fluorescent siNA complexed with lipid. Cells in 6-well dishes are incubated with siNA for 24 hours, rinsed with PBS and fixed in 2% paraformaldehyde for 15 minutes at room temperature. Uptake of siNA is visualized using a fluorescent microscope.

#### Tagman and Lightcycler quantification of mRNA

Total RNA is prepared from cells following siNA delivery, for example, using Qiagen RNA purification kits for 6-well or Rneasy extraction kits for 96-well assays. For

Taqman analysis, dual-labeled probes are synthesized with the reporter dye, FAM or JOE, covalently linked at the 5'-end and the quencher dye TAMRA conjugated to the 3'-end. One-step RT-PCR amplifications are performed on, for example, an ABI PRISM 7700 Sequence Detector using 50 µl reactions consisting of 10 µl total RNA, 100 nM forward primer, 900 nM reverse primer, 100 nM probe, 1X TaqMan PCR reaction buffer (PE-Applied Biosystems), 5.5 mM MgCl<sub>2</sub>, 300 μM each dATP, dCTP, dGTP, and dTTP, 10U RNase Inhibitor (Promega), 1.25U AmpliTaq Gold (PE-Applied Biosystems) and 10U M-MLV Reverse Transcriptase (Promega). The thermal cycling conditions can consist of 30 min at 48°C, 10 min at 95°C, followed by 40 cycles of 15 sec at 95°C and 1 min at 60°C. Quantitation of mRNA levels is determined relative to standards generated from serially diluted total cellular RNA (300, 100, 33, 11 ng/rxn) and normalizing to \(\beta\)-actin or GAPDH mRNA in parallel TaqMan reactions. For each gene of interest an upper and lower primer and a fluorescently labeled probe are designed. Real time incorporation of SYBR Green I dye into a specific PCR product can be measured in glass capillary tubes using a lightcyler. A standard curve is generated for each primer pair using control cRNA. Values are represented as relative expression to GAPDH in each sample.

## Western blotting

5

10

15

20

25

30

Nuclear extracts can be prepared using a standard micro preparation technique (see for example Andrews and Faller, 1991, *Nucleic Acids Research*, 19, 2499). Protein extracts from supernatants are prepared, for example using TCA precipitation. An equal volume of 20% TCA is added to the cell supernatant, incubated on ice for 1 hour and pelleted by centrifugation for 5 minutes. Pellets are washed in acetone, dried and resuspended in water. Cellular protein extracts are run on a 10% Bis-Tris NuPage (nuclear extracts) or 4-12% Tris-Glycine (supernatant extracts) polyacrylamide gel and transferred onto nitro-cellulose membranes. Non-specific binding can be blocked by incubation, for example, with 5% non-fat milk for 1 hour followed by primary antibody for 16 hour at 4°C. Following washes, the secondary antibody is applied, for example (1:10,000 dilution) for 1 hour at room temperature and the signal detected with SuperSignal reagent (Pierce).

# Example 11: Animal Models

Various animal models can be used to screen siNA constructs in vivo as are known in the art, for example those animal models that are used to evaluate other nucleic acid technologies such as enzymatic nucleic acid molecules (ribozymes) and/or antisense. Such animal models are used to test the efficacy of siNA molecules described herein. In a non-limiting example, siNA molecules that are designed as anti-angiogenic agents can There are several animal models in which the antibe screened animal models. angiogenesis effect of nucleic acids of the present invention, such as siNA, directed against genes associated with angiogenesis and/or metastais, such as VEGFR (e.g., VEGFR1, VEGFR2, and VEGFR3) genes. Typically a corneal model has been used to study angiogenesis in rat and rabbit since recruitment of vessels can easily be followed in this normally avascular tissue (Pandey et al., 1995 Science 268: 567-569). In these models, a small Teflon or Hydron disk pretreated with an angiogenesis factor (e.g. bFGF or VEGF) is inserted into a pocket surgically created in the cornea. Angiogenesis is monitored 3 to 5 days later. siNA molecules directed against VEGFR mRNAs are delivered in the disk as well, or dropwise to the eye over the time course of the experiment. In another eye model, hypoxia has been shown to cause both increased expression of VEGF and neovascularization in the retina (Pierce et al., 1995 Proc. Natl. Acad. Sci. USA. 92: 905-909; Shweiki et al., 1992 J. Clin. Invest. 91: 2235-2243).

Several animal models exist for screening of anti-angiogenic agents. These include corneal vessel formation following corneal injury (Burger et al., 1985 Cornea 4: 35-41; Lepri, et al., 1994 J. Ocular Pharmacol. 10: 273-280; Ormerod et al., 1990 Am. J. Pathol. 137: 1243-1252) or intracorneal growth factor implant (Grant et al., 1993 Diabetologia 36: 282-291; Pandey et al. 1995 supra; Zieche et al., 1992 Lab. Invest. 67: 711-715), vessel growth into Matrigel matrix containing growth factors (Passaniti et al., 1992 supra), female reproductive organ neovascularization following hormonal manipulation (Shweiki et al., 1993 Clin. Invest. 91: 2235-2243), several models involving inhibition of tumor growth in highly vascularized solid tumors (O'Reilly et al., 1994 Cell 79: 315-328; Senger et al., 1993 Cancer and Metas. Rev. 12: 303-324; Takahasi et al., 1994 Cancer Res. 54: 4233-4237; Kim et al., 1993 supra), and transient hypoxia-induced neovascularization in the mouse retina (Pierce et al., 1995 Proc. Natl. Acad. Sci. USA. 92: 905-909).gene

5

10

15

20

25

The cornea model, described in Pandey et al. *supra*, is the most common and well characterized anti-angiogenic agent efficacy screening model. This model involves an avascular tissue into which vessels are recruited by a stimulating agent (growth factor, thermal or alkalai burn, endotoxin). The corneal model would utilize the intrastromal corneal implantation of a Teflon pellet soaked in a VEGF-Hydron solution to recruit blood vessels toward the pellet which can be quantitated using standard microscopic and image analysis techniques. To evaluate their anti-angiogenic efficacy, ribozymes are applied topically to the eye or bound within Hydron on the Teflon pellet itself. This avascular cornea as well as the Matrigel model provide for low background assays. While the corneal model has been performed extensively in the rabbit, studies in the rat have also been conducted.

The mouse model (Passaniti et al., *supra*) is a non-tissue model which utilizes Matrigel, an extract of basement membrane (Kleinman et al., 1986) or Millipore<sup>®</sup> filter disk, which can be impregnated with growth factors and anti-angiogenic agents in a liquid form prior to injection. Upon subcutaneous administration at body temperature, the Matrigel or Millipore<sup>®</sup> filter disk forms a solid implant. VEGF embedded in the Matrigel or Millipore<sup>®</sup> filter disk is used to recruit vessels within the matrix of the Matrigel or Millipore<sup>®</sup> filter disk which can be processed histologically for endothelial cell specific vWF (factor VIII antigen) immunohistochemistry, Trichrome-Masson stain, or hemoglobin content. Like the cornea, the Matrigel or Millipore<sup>®</sup> filter disk are avascular; however, it is not tissue. In the Matrigel or Millipore<sup>®</sup> filter disk model, siNA molecules are administered within the matrix of the Matrigel or Millipore<sup>®</sup> filter disk to test their anti-angiogenic efficacy. Thus, delivery issues in this model, as with delivery of siNA molecules by Hydron- coated Teflon pellets in the rat comea model, may be less problematic due to the homogeneous presence of the siNA within the respective matrix.

The Lewis lung carcinoma and B-16 murine melanoma models are well accepted models of primary and metastatic cancer and are used for initial screening of anti-cancer agents. These murine models are not dependent upon the use of immunodeficient mice, are relatively inexpensive, and minimize housing concerns. Both the Lewis lung and B-16 melanoma models involve subcutaneous implantation of approximately  $10^6$  tumor cells from metastatically aggressive tumor cell lines (Lewis lung lines 3LL or D122, LLc-

5

10

15

20

25

LN7; B-16-BL6 melanoma) in C57BL/6J mice. Alternatively, the Lewis lung model can be produced by the surgical implantation of tumor spheres (approximately 0.8 mm in diameter). Metastasis also may be modeled by injecting the tumor cells directly *i.v.*. In the Lewis lung model, microscopic metastases can be observed approximately 14 days following implantation with quantifiable macroscopic metastatic tumors developing within 21-25 days. The B-16 melanoma exhibits a similar time course with tumor neovascularization beginning 4 days following implantation. Since both primary and metastatic tumors exist in these models after 21-25 days in the same animal, multiple measurements can be taken as indices of efficacy. Primary tumor volume and growth latency as well as the number of micro- and macroscopic metastatic lung foci or number of animals exhibiting metastases can be quantitated. The percent increase in lifespan can also be measured. Thus, these models provide suitable primary efficacy assays for screening systemically administered siNA molecules and siNA formulations.

In the Lewis lung and B-16 melanoma models, systemic pharmacotherapy with a wide variety of agents usually begins 1-7 days following tumor implantation/inoculation with either continuous or multiple administration regimens. Concurrent pharmacokinetic studies can be performed to determine whether sufficient tissue levels of siNA can be achieved for pharmacodynamic effect to be expected. Furthermore, primary tumors and secondary lung metastases can be removed and subjected to a variety of *in vitro* studies (*i.e.* target RNA reduction).

In utilizing these models to assess siNA activity, VEGFR1, VEGFR2, and/or VEGFR3 protein levels can be measured clinically or experimentally by FACS analysis. VEGFR1, VEGFR2, and/or VEGFR3 encoded mRNA levels will be assessed by Northern analysis, RNase-protection, primer extension analysis and/or quantitative RT-PCR. siNA molecules that block VEGFR1, VEGFR2, and/or VEGFR3 protein encoding mRNAs and therefore result in decreased levels of VEGFR1, VEGFR2, and/or VEGFR3 activity by more than 20% *in vitro* can be thus identified.

### Example 12: siNA-mediated inhibition of angiogenesis in vivo

The purpose of this study was to assess the anti-angiogenic activity of siNA targeted against VEGFR1 in the rat cornea model of VEGF induced angiogenesis (see above). These siNA molecules have matched inverted controls which are inactive since

5

10

15

20

25

they are not able to interact with the RNA target. The siNA molecules and VEGF were co-delivered using the filter disk method: Nitrocellulose filter disks (Millipore<sup>®</sup>) of 0.057 diameter were immersed in appropriate solutions and were surgically implanted in rat cornea as described by Pandey *et al.*, *supra*.

The stimulus for angiogenesis in this study was the treatment of the filter disk with 30 µM VEGF which is implanted within the cornea's stroma. This dose yields reproducible neovascularization stemming from the pericorneal vascular plexus growing toward the disk in a dose-response study 5 days following implant. Filter disks treated only with the vehicle for VEGF show no angiogenic response. The siNA were coadminstered with VEGF on a disk in two different siNA concentrations. One concern with the simultaneous administration is that the siNA would not be able to inhibit angiogenesis since VEGF receptors can be stimulated. However, Applicant has observed that in low VEGF doses, the neovascular response reverts to normal, suggesting that the VEGF stimulus is essential for maintaining the angiogenic response. Blocking the production of VEGF receptors using simultaneous administration of anti-VEGF-R mRNA siNA could attenuate the normal neovascularization induced by the filter disk treated with VEGF.

## Materials and Methods:

Test Compounds and Controls

20

5

10

15

R&D Systems VEGF, carrier free at 75  $\mu$ M in 82 mM Tris-Cl, pH 6.9 siNA, 1.67  $\mu$ G/ $\mu$ L, SITE 2340 (SEQ ID NO: 2; SEQ ID NO: 6) sense/antisense siNA, 1.67  $\mu$ G/ $\mu$ L, INVERTED CONTROL FOR SITE 2340 (SEQ ID NO: 19; SEQ ID NO: 20) sense/antisense

25

siNA 1.67  $\mu$ g/ $\mu$ L, Site 2340 (SEQ ID NO: 419; SEQ ID NO: 420) sense/antisense

## Animals

Harlan Sprague-Dawley Rats, Approximately 225-250g
45 males, 5 animals per group.

#### Husbandry

Animals are housed in groups of two. Feed, water, temperature and humidity are determined according to Pharmacology Testing Facility performance standards (SOP's) which are in accordance with the 1996 Guide for the Care and Use of Laboratory Animals (NRC). Animals are acclimated to the facility for at least 7 days prior to experimentation. During this time, animals are observed for overall health and sentinels will be bled for baseline serology.

## Experimental Groups

10

5

Each solution (VEGF and siNAs) was prepared as a 1X solution for final concentrations shown in the experimental groups described in Table III.

## siNA Annealing Conditions

15

siNA sense and antisense strands are annealed for 1 minute in  $H_2O$  at 1.67mg/mL/strand followed by a 1 hour incubation at 37°C producing 3.34 mg/mL of duplexed siNA. For the  $20\mu g$ /eye treatment, 6  $\mu$ Ls of the 3.34 mg/mL duplex is injected into the eye (see below). The 3.34 mg/mL duplex siNA can then be serially diluted for dose response assays.

20

25

## Preparation of VEGF Filter Disk

For corneal implantation, 0.57 mm diameter nitrocellulose disks, prepared from 0.45  $\mu$ m pore diameter nitrocellulose filter membranes (Millipore Corporation), were soaked for 30 min in 1  $\mu$ L of 75  $\mu$ M VEGF in 82 mM Tris HCl (pH 6.9) in covered petri dishes on ice. Filter disks soaked only with the vehicle for VEGF (83 mM Tris-Cl pH 6.9) elicit no angiogenic response.

#### Corneal surgery

The rat corneal model used in this study was a modified from Koch et al. Supra and Pandey et al., supra. Briefly, corneas were irrigated with 0.5% povidone iodine solution followed by normal saline and two drops of 2% lidocaine. Under a dissecting microscope (Leica MZ-6), a stromal pocket was created and a presoaked filter disk (see above) was inserted into the pocket such that its edge was 1 mm from the corneal limbus.

## Intraconjunctival injection of test solutions

5

10

15

20

25

Immediately after disk insertion, the tip of a 40-50 µm OD injector (constructed in our laboratory) was inserted within the conjunctival tissue 1 mm away from the edge of the corneal limbus that was directly adjacent to the VEGF-soaked filter disk. Six hundred nanoliters of test solution (siNA, inverted control or sterile water vehicle) were dispensed at a rate of 1.2 µL/min using a syringe pump (Kd Scientific). The injector was then removed, serially rinsed in 70% ethanol and sterile water and immersed in sterile water between each injection. Once the test solution was injected, closure of the eyelid was maintained using microaneurism clips until the animal began to recover gross motor activity. Following treatment, animals were warmed on a heating pad at 37°C.

## Quantitation of angiogenic response

Five days after disk implantation, animals were euthanized following im administration of 0.4 mg/kg atropine and corneas were digitally imaged. The neovascular surface area (NSA, expressed in pixels) was measured *postmortem* from blood-filled corneal vessels using computerized morphometry (Image Pro Plus, Media Cybernetics, v2.0). The individual mean NSA was determined in triplicate from three regions of identical size in the area of maximal neovascularization between the filter disk and the limbus. The number of pixels corresponding to the blood-filled corneal vessels in these regions was summated to produce an index of NSA. A group mean NSA was then calculated. Data from each treatment group were normalized to VEGF/siNA vehicle-treated control NSA and finally expressed as percent inhibition of VEGF-induced angiogenesis.

#### 30 Statistics

After determining the normality of treatment group means, group mean percent inhibition of VEGF-induced angiogenesis was subjected to a one-way analysis of variance. This was followed by two post-hoc tests for significance including Dunnett's (comparison to VEGF control) and Tukey-Kramer (all other group mean comparisons) at alpha = 0.05. Statistical analyses were performed using JMP v.3.1.6 (SAS Institute).

Results are graphically represented in Figure 23. As shown in Figure 23, VEGFR1 site 4229 active siNA at three concentrations were effective at inhibiting angiogenesis compared to the inverted siNA control and the VEGF control. A chemically modified version of the VEGFR1 site 4229 active siNA comprising a sense strand having 2'-deoxy-2'-fluoro pyrimidines and ribo purines with 5' and 3' terminal inverted deoxyabasic residues (SEQ ID NO: 419) and an antisense strand having having 2'-deoxy-2'-fluoro pyrimidines and ribo purines with a terminal 3'-phosphorothioate internucleotide linkage (SEQ ID NO: 420), showed similar inhibition. This result shows siNA molecules of differing chemically modified composition of the invention are capable of significantly inhibiting angiogenesis in vivo.

## Example 13: RNAi mediated inhibition of EGFR (HER1) RNA expression

siNA constructs (Table I) were tested for efficacy in reducing EGFR (HER1) RNA expression in A549 cells. A549 cells were plated approximately 24h before transfection in 96-well plates at 5,000-7,500 cells/well, 100 µl/well, such that at the time of transfection cells are 70-90% confluent. For transfection, annealed siNAs were mixed with the transfection reagent (Lipofectamine 2000, Invitrogen) in a volume of 50 µl/well and incubated for 20 min. at room temperature. The siNA transfection mixtures were added to cells to give a final siNA concentration of 25 nM in a volume of 150 µl. Each siNA transfection mixture was added to 3 wells for triplicate siNA treatments. Cells were incubated at 37°C for 24h in the continued presence of the siNA transfection mixture. At 24h, RNA was prepared from each well of treated cells. The supernatants with the transfection mixtures were first removed and discarded, then the cells wre lysed and RNA prepared from each well. Target gene expression following treatment was evaluated by RT-PCR for the target gene and for a control gene (36B4, an RNA polymerase subunit) The triplicate data were averaged and the standard deviations for normalization. determined for each treatment. Normalized data were graphed and the percent reduction

5

10

15

20

25

of target mRNA by active siNAs in comparison to their respective inverted control siNAs was determined.

Results of this study are shown in **Figure 25**. A siNA construct comprising ribonucleotides and 3'-terminal dithymidine caps (RPI#30988/31064) was compared to a chemically modified siNA construct comprising 2'-deoxy-2'-fluoro pyrimidine nucleotides and purine ribonucleotides in which the sense strand of the siNA is further modified with 5' and 3'-terminal inverted deoxyabasic caps and the antisense strand comprises a 3'-terminal phosphorothioate internucleotide linkage (RPI#31300/31301), which was also compared to a matched chemistry inverted control (RPI#31312/31313). In addition, the siNA constructs were also compared to untreated cells, cells transfected with lipid and scrambled siNA constructs (Scram1 and Scram2), and cells transfected with lipid alone (transfection control). As shown in the figure, both siNA constructs significantly reduce EGFR RNA expression. Additional stabilization chemistries as described in **Table IV** are similarly assayed for activity.

# Example 14: RNAi mediated inhibition of PKC-alpha RNA expression

siNA constructs (Table I) are tested for efficacy in reducing PKC-alpha RNA expression in, for example in A549 cells. Cells are plated approximately 24h before transfection in 96-well plates at 5,000-7,500 cells/well, 100 µl/well, such that at the time of transfection cells are 70-90% confluent. For transfection, annealed siNAs are mixed with the transfection reagent (Lipofectamine 2000, Invitrogen) in a volume of 50 µl/well and incubated for 20 min. at room temperature. The siNA transfection mixtures are added to cells to give a final siNA concentration of 25 nM in a volume of 150 µl. Each siNA transfection mixture is added to 3 wells for triplicate siNA treatments. Cells are incubated at 37° for 24h in the continued presence of the siNA transfection mixture. At 24h, RNA is prepared from each well of treated cells. The supernatants with the transfection mixtures are first removed and discarded, then the cells are lysed and RNA prepared from each well. Target gene expression following treatment is evaluated by RT-PCR for the target gene and for a control gene (36B4, an RNA polymerase subunit) for normalization. The triplicate data is averaged and the standard deviations determined for each treatment. Normalized data are graphed and the percent reduction of target mRNA by active siNAs in comparison to their respective inverted control siNAs was determined.

5

10

15

20

25

In a non-limiting example, siNA constructs were screened for activity (see Figure 26) and compared to untreated cells, scrambled siNA control constructs (Scram1 and Scram2), and cells transfected with lipid alone (transfection control). As shown in Figure 26, the siNA constructs significantly reduce PKC-alpha RNA expression. Leads generated from such a screen are then further assayed. In a non-limiting example, siNA constructs comprising ribonucleotides and 3'-terminal dithymidine caps are assayed along with a chemically modified siNA construct comprising 2'-deoxy-2'-fluoro pyrimidine nucleotides and purine ribonucleotides, in which the sense strand of the siNA is further modified with 5' and 3'-terminal inverted deoxyabasic caps and the antisense strand comprises a 3'-terminal phosphorothioate internucleotide linkage. Additional stabilization chemistries as described in Table IV are similarly assayed for activity. These siNA constructs are compared to appropriate matched chemistry inverted controls. In addition, the siNA constructs are also compared to untreated cells, cells transfected with lipid and scrambled siNA constructs, and cells transfected with lipid alone (transfection control).

## Example 15: RNAi mediated inhibition of Myc RNA expression

siNA constructs (Table I) were tested for efficacy in reducing Myc (c-Myc) RNA expression in 293T cells. 293T cells were plated approximately 24h before transfection in 96-well plates at 5,000-7,500 cells/well, 100 µl/well, such that at the time of transfection cells were 70-90% confluent. For transfection, annealed siNAs were mixed with the transfection reagent (Lipofectamine 2000, Invitrogen) in a volume of 50 µl/well and incubated for 20 min. at room temperature. The siNA transfection mixtures were added to cells to give a final siNA concentration of 25 nM in a volume of 150 µl. Each siNA transfection mixture was added to 3 wells for triplicate siNA treatments. Cells were incubated at 37°C for 24h in the continued presence of the siNA transfection mixture. At 24h, RNA was prepared from each well of treated cells. The supernatants with the transfection mixtures were first removed and discarded, then the cells were lysed and RNA prepared from each well. Target gene expression following treatment was evaluated by RT-PCR for the target gene and for a control gene (36B4, an RNA polymerase subunit) for normalization. The triplicate data were averaged and the standard deviations determined for each treatment. Normalized data were graphed and

5

10

15

20

25

the percent reduction of target mRNA by active siNAs in comparison to their respective inverted control siNAs was determined.

Results of this study are shown in **Figure 27**. A screen of siNA constructs was compared to untreated cells, scrambled siNA control constructs (Scram1 and Scram2), and cells transfected with lipid alone (transfection control). As shown in the figure, three of the siNA constructs (RPI 30993/31069; RPI 30995/31071; and RPI 30996/31072) significantly reduce c-Myc RNA expression. Additional stabilization chemistries as described in **Table IV** are similarly assayed for activity.

# Example 16: RNAi mediated inhibition of BCL2 RNA expression

siNA constructs (Table I) are tested for efficacy in reducing BCL2 RNA expression in, for example, A549 cells. Cells are plated approximately 24h before transfection in 96well plates at 5,000-7,500 cells/well, 100 µl/well, such that at the time of transfection cells are 70-90% confluent. For transfection, annealed siNAs are mixed with the transfection reagent (Lipofectamine 2000, Invitrogen) in a volume of 50 µl/well and incubated for 20 min. at room temperature. The siNA transfection mixtures are added to cells to give a final siNA concentration of 25 nM in a volume of 150 µl. Each siNA transfection mixture is added to 3 wells for triplicate siNA treatments. Cells are incubated at 37° for 24h in the continued presence of the siNA transfection mixture. At 24h, RNA is prepared from each well of treated cells. The supernatants with the transfection mixtures are first removed and discarded, then the cells are lysed and RNA prepared from each well. Target gene expression following treatment is evaluated by RT-PCR for the target gene and for a control gene (36B4, an RNA polymerase subunit) for normalization. The triplicate data is averaged and the standard deviations determined for each treatment. Normalized data are graphed and the percent reduction of target mRNA by active siNAs in comparison to their respective inverted control siNAs is determined.

In a non-limiting example, A549 cells were transfected with 0.25 ug/well of lipid complexed with 25 nM siNA. A siNA construct comprising ribonucleotides and 3'-terminal dithymidine caps (RPI#30998/31074) was tested along with a chemically modified siNA construct comprising 2'-deoxy-2'-fluoro pyrimidine nucleotides and purine ribonucleotides in which the sense strand of the siNA is further modified with 5' and 3'-terminal inverted deoxyabasic caps and the antisense strand comprises a 3'-

5

10

15

20

25

terminal phosphorothioate internucleotide linkage (RPI#31368/31369), which was also compared to a matched chemistry inverted control (RPI#31370/31371) and a chemically modified siNA construct comprising 2'-deoxy-2'-fluoro pyrimidine and 2'-deoxy-2'-fluoro purine nucleotides in which the sense strand of the siNA is further modified with 5' and 3'-terminal inverted deoxyabasic caps and the antisense strand comprises a 3'-terminal phosphorothioate internucleotide linkage (RPI#31372/31373) which was also compared to a matched chemistry inverted control (RPI#31374/31375). In addition, the siNA constructs were also compared to untreated cells, cells transfected with lipid and scrambled siNA constructs (Scram1 and Scram2), and cells transfected with lipid alone (transfection control). As shown in Figure 28, the siNA constructs significantly reduce BCL2 RNA expression compared to scrambled, untreated, and transfection controls. Additional stabilization chemistries as described in Table IV are similarly assayed for activity.

## Example 17: RNAi mediated inhibition of CHK-1 RNA expression

siNA constructs (Table I) were tested for efficacy in reducing CHK-1 RNA expression in A549 cells. A549 cells were plated approximately 24h before transfection in 96-well plates at 5,000-7,500 cells/well, 100 µl/well, such that at the time of transfection cells are 70-90% confluent. For transfection, annealed siNAs were mixed with the transfection reagent (Lipofectamine 2000, Invitrogen) in a volume of 50 μl/well and incubated for 20 min. at room temperature. The siNA transfection mixtures were added to cells to give a final siNA concentration of 25 nM in a volume of 150 µl. Each siNA transfection mixture was added to 3 wells for triplicate siNA treatments. Cells were incubated at 37° for 24h in the continued presence of the siNA transfection mixture. At 24h, RNA was prepared from each well of treated cells. The supernatants with the transfection mixtures were first removed and discarded, then the cells were lysed and RNA prepared from each well. Target gene expression following treatment was evaluated by RT-PCR for the target gene and for a control gene (36B4, an RNA polymerase subunit) for normalization. The triplicate data were averaged and the standard deviations determined for each treatment. Normalized data were graphed and the percent reduction of target mRNA by active siNAs in comparison to their respective inverted control siNAs was determined.

5

10

15

20

25

Results of this study are shown in **Figure 29**. A siNA construct comprising ribonucleotides and 3'-terminal dithymidine caps (RPI#31003/31079) and a chemically modified siNA construct comprising 2'-deoxy-2'-fluoro pyrimidine nucleotides and purine ribonucleotides in which the sense strand of the siNA is further modified with 5' and 3'-terminal inverted deoxyabasic caps and in which the antisense strand comprises a 3'-terminal phosphorothioate internucleotide linkage (RPI#31302/31303), were compared to a matched chemistry inverted control (RPI#31314/31325). In addition, the siNA constructs were also compared to untreated cells, cells transfected with lipid and scrambled siNA constructs (Scram1 and Scram2), and cells transfected with lipid alone (transfection control). As shown in the figure, both siNA constructs significantly reduce CHK-1 RNA expression compared to appropriate controls. Additional stabilization chemistries as described in **Table IV** are similarly assayed for activity.

## Example 18: RNAi mediated inhibition of BACE RNA expression

siNA constructs (Table I) are tested for efficacy in reducing BACE RNA expression in, for example in A549 cells. Cells are plated approximately 24h before transfection in 96well plates at 5,000-7,500 cells/well, 100 µl/well, such that at the time of transfection cells are 70-90% confluent. For transfection, annealed siNAs are mixed with the transfection reagent (Lipofectamine 2000, Invitrogen) in a volume of 50 µl/well and incubated for 20 min. at room temperature. The siNA transfection mixtures are added to cells to give a final siNA concentration of 25 nM in a volume of 150 µl. Each siNA transfection mixture is added to 3 wells for triplicate siNA treatments. Cells are incubated at 37°C for 24h in the continued presence of the siNA transfection mixture. At 24h, RNA is prepared from each well of treated cells. The supernatants with the transfection mixtures are first removed and discarded, then the cells are lysed and RNA prepared from each well. Target gene expression following treatment is evaluated by RT-PCR for the target gene and for a control gene (36B4, an RNA polymerase subunit) for normalization. The triplicate data is averaged and the standard deviations determined for each treatment. Normalized data are graphed and the percent reduction of target mRNA by active siNAs in comparison to their respective inverted control siNAs was determined.

In a non-limiting example, siNA constructs were screened for activity (see Figure 30) and compared to untreated cells, scrambled siNA control constructs (Scram1 and

5

10

15

20

25

Scram2), and cells transfected with lipid alone (transfection control). As shown in Figure 30, the siNA constructs significantly reduce BACE RNA expression. Leads generated from such a screen are then further assayed. In a non-limiting example, siNA constructs comprising ribonucleotides and 3'-terminal dithymidine caps are assayed along with a chemically modified siNA construct comprising 2'-deoxy-2'-fluoro pyrimidine nucleotides and purine ribonucleotides, in which the sense strand of the siNA is further modified with 5' and 3'-terminal inverted deoxyabasic caps and the antisense strand comprises a 3'-terminal phosphorothioate internucleotide linkage. Additional stabilization chemistries as described in Table IV are similarly assayed for activity. These siNA constructs are compared to appropriate matched chemistry inverted controls. In addition, the siNA constructs are also compared to untreated cells, cells transfected with lipid and scrambled siNA constructs, and cells transfected with lipid alone (transfection control).

## Example 19: RNAi mediated inhibition of cyclin D1 RNA expression

siNA constructs (Table I) were tested for efficacy in reducing cyclin D1 RNA expression in A549 cells. A549 cells were plated approximately 24h before transfection in 96-well plates at 5,000-7,500 cells/well, 100 µl/well, such that at the time of transfection cells are 70-90% confluent. For transfection, annealed siNAs were mixed with the transfection reagent (Lipofectamine 2000, Invitrogen) in a volume of 50 µl/well and incubated for 20 min. at room temperature. The siNA transfection mixtures were added to cells to give a final siNA concentration of 25 nM in a volume of 150 µl. Each siNA transfection mixture was added to 3 wells for triplicate siNA treatments. Cells were incubated at 37° for 24h in the continued presence of the siNA transfection mixture. At 24h, RNA was prepared from each well of treated cells. The supernatants with the transfection mixtures were first removed and discarded, then the cells were lysed and RNA prepared from each well. Target gene expression following treatment was evaluated by RT-PCR for the target gene and for a control gene (36B4, an RNA polymerase subunit) for normalization. The triplicate data were averaged and the standard deviations determined for each treatment. Normalized data were graphed and the percent reduction of target mRNA by active siNAs in comparison to their respective inverted control siNAs was determined.

5

10

15

20

25

Results of this study are shown in Figure 31. A siNA construct comprising ribonucleotides and 3'-terminal dithymidine caps (RPI#30988/31064) was assayed along with a chemically modified siNA construct comprising 2'-deoxy-2'-fluoro pyrimidine nucleotides and purine ribonucleotides in which the sense strand of the siNA is further modified with 5' and 3'-terminal inverted deoxyabasic caps and the antisense strand comprises a 3'-terminal phosphorothioate internucleotide linkage (RPI#31300/3130), which was also compared to a matched chemistry inverted control (RPI#31312/31313). In addition, the siNA constructs were also compared to untreated cells, cells transfected with lipid and scrambled siNA constructs (Scram1 and Scram2), and cells transfected with lipid alone (transfection control). As shown in the figure, both siNA constructs significantly reduce cyclin D1 RNA expression. Additional stabilization chemistries as described in Table IV are similarly assayed for activity.

## Example 20: RNAi mediated inhibition of PTP-1B RNA expression

siNA constructs (Table I) were tested for efficacy in reducing PTP-1B RNA expression in A549 cells. A549 cells were plated approximately 24h before transfection in 96-well plates at 5,000-7,500 cells/well, 100 µl/well, such that at the time of transfection cells are 70-90% confluent. For transfection, annealed siNAs were mixed with the transfection reagent (Lipofectamine 2000, Invitrogen) in a volume of 50 µl/well and incubated for 20 min. at room temperature. The siNA transfection mixtures were added to cells to give a final siNA concentration of 25 nM in a volume of 150 µl. Each siNA transfection mixture was added to 3 wells for triplicate siNA treatments. Cells were incubated at 37° for 24h in the continued presence of the siNA transfection mixture. At 24h, RNA was prepared from each well of treated cells. The supernatants with the transfection mixtures were first removed and discarded, then the cells were lysed and RNA prepared from each well. Target gene expression following treatment was evaluated by RT-PCR for the target gene and for a control gene (36B4, an RNA polymerase subunit) for normalization. The triplicate data were averaged and the standard deviations determined for each treatment. Normalized data were graphed and the percent reduction of target mRNA by active siNAs in comparison to their respective inverted control siNAs was determined.

5

10

15

20

25

Results of this study are shown in Figure 32. A siNA construct comprising ribonucleotides and 3'-terminal dithymidine caps (RPI#31018/31094) was assayed along with a chemically modified siNA construct comprising 2'-deoxy-2'-fluoro pyrimidine nucleotides and purine ribonucleotides in which the sense strand of the siNA is further modified with 5' and 3'-terminal inverted deoxyabasic caps and the antisense strand comprises a 3'-terminal phosphorothioate internucleotide linkage (RPI#31306/31307), which was also compared to a matched chemistry inverted control (RPI#31318/31319). In addition, the siNA constructs were also compared to untreated cells, cells transfected with lipid and scrambled siNA constructs (Scram1 and Scram2), and cells transfected with lipid alone (transfection control). As shown in the figure, both siNA constructs significantly reduce PTP-1B RNA expression. Additional stabilization chemistries as described in Table IV are similarly assayed for activity.

## Example 21: RNAi mediated inhibition of ERG2 RNA expression

siNA constructs (Table I) are tested for efficacy in reducing ERG2 RNA expression in, for example in DLD1 cells. Cells are plated approximately 24h before transfection in 96-well plates at 5,000-7,500 cells/well, 100 µl/well, such that at the time of transfection cells are 70-90% confluent. For transfection, annealed siNAs are mixed with the transfection reagent (Lipofectamine 2000, Invitrogen) in a volume of 50 µl/well and incubated for 20 min. at room temperature. The siNA transfection mixtures are added to cells to give a final siNA concentration of 25 nM in a volume of 150  $\mu$ l. Each siNA transfection mixture is added to 3 wells for triplicate siNA treatments. Cells are incubated at 37° for 24h in the continued presence of the siNA transfection mixture. At 24h, RNA is prepared from each well of treated cells. The supernatants with the transfection mixtures are first removed and discarded, then the cells are lysed and RNA prepared from each well. Target gene expression following treatment is evaluated by RT-PCR for the target gene and for a control gene (36B4, an RNA polymerase subunit) for normalization. The triplicate data is averaged and the standard deviations determined for each treatment. Normalized data are graphed and the percent reduction of target mRNA by active siNAs in comparison to their respective inverted control siNAs was determined.

In a non-limiting example, siNA constructs were screened for activity (see Figure 33) and compared to untreated cells, scrambled siNA control constructs (Scram1 and

5

10

15

20

25

Scram2), and cells transfected with lipid alone (transfection control). As shown in Figure 33, the siNA constructs significantly reduce of ERG2 RNA expression. Leads generated from such a screen are then further assayed. In a non-limiting example, siNA constructs comprising ribonucleotides and 3'-terminal dithymidine caps are assayed along with a chemically modified siNA construct comprising 2'-deoxy-2'-fluoro pyrimidine nucleotides and purine ribonucleotides, in which the sense strand of the siNA is further modified with 5' and 3'-terminal inverted deoxyabasic caps and the antisense strand comprises a 3'-terminal phosphorothioate internucleotide linkage. Additional stabilization chemistries as described in Table IV are similarly assayed for activity. These siNA constructs are compared to appropriate matched chemistry inverted controls. In addition, the siNA constructs are also compared to untreated cells, cells transfected with lipid and scrambled siNA constructs, and cells transfected with lipid alone (transfection control). Additional stabilization chemistries as described in Table IV are similarly assayed for activity.

## 15 Example 22: RNAi mediated inhibition of PCNA RNA expression

siNA constructs (Table I) were tested for efficacy in reducing PCNA RNA expression in A549 cells. A549 cells were plated approximately 24h before transfection in 96-well plates at 5,000-7,500 cells/well, 100  $\mu$ l/well, such that at the time of transfection cells are 70-90% confluent. For transfection, annealed siNAs were mixed with the transfection reagent (Lipofectamine 2000, Invitrogen) in a volume of 50 µl/well and incubated for 20 min. at room temperature. The siNA transfection mixtures were added to cells to give a final siNA concentration of 25 nM in a volume of 150 µl. Each siNA transfection mixture was added to 3 wells for triplicate siNA treatments. Cells were incubated at 37° for 24h in the continued presence of the siNA transfection mixture. At 24h, RNA was prepared from each well of treated cells. The supernatants with the transfection mixtures were first removed and discarded, then the cells were lysed and RNA prepared from each well. Target gene expression following treatment was evaluated by RT-PCR for the target gene and for a control gene (36B4, an RNA polymerase subunit) for normalization. The triplicate data were averaged and the standard deviations determined for each treatment. Normalized data were graphed and the percent reduction of target mRNA by active siNAs in comparison to their respective inverted control siNAs was determined.

5

10

20

25

Results of this study are shown in Figure 34. A siNA construct comprising ribonucleotides and 3'-terminal dithymidine caps (RPI#31035/31111) was assayed along with a chemically modified siNA construct comprising 2'-deoxy-2'-fluoro pyrimidine nucleotides and purine ribonucleotides in which the sense strand of the siNA is further modified with 5' and 3'-terminal inverted deoxyabasic caps and the antisense strand comprises a 3'-terminal phosphorothioate internucleotide linkage (RPI#31310/31311), which was also compared to a matched chemistry inverted control (RPI#31322/31323). In addition, the siNA constructs were also compared to untreated cells, cells transfected with lipid and scrambled siNA constructs (Scram1 and Scram2), and cells transfected with lipid alone (transfection control). As shown in the figure, both siNA constructs significant reduce PCNA RNA expression. Additional stabilization chemistries as described in Table IV are similarly assayed for activity.

#### Example 23: Indications

5

10

15

20

25

30

The siNA molecules of the invention can be used to treat a variety of diseases and conditions through modulation of gene expression. Using the methods described herein, chemically modified siNA molecules can be designed to modulate the expression any number of target genes, including but not limited to genes associated with cancer, metabolic diseases, infectious diseases such as viral, bacterial or fungal infections, neurologic diseases, musculoskeletal diseases, diseases of the immune system, diseases associated with signaling pathways and cellular messengers, and diseases associated with transport systems including molecular pumps and channels.

Non-limiting examples of various viral genes that can be targeted using siRNA molecules of the invention include Hepatitis C Virus (HCV, for example Genbank Accession Nos: D11168, D50483.1, L38318 and S82227), Hepatitis B Virus (HBV, for example GenBank Accession No. AF100308.1), Human Immunodeficiency Virus type 1 (HIV-1, for example GenBank Accession No. U51188), Human Immunodeficiency Virus type 2 (HIV-2, for example GenBank Accession No. X60667), West Nile Virus (WNV for example GenBank accession No. NC\_001563), cytomegalovirus (CMV for example GenBank Accession No. NC\_001347), respiratory syncytial virus (RSV for example GenBank Accession No. NC\_001781), influenza virus (for example GenBank Accession numbers:

D00239, X02316, X01087, L24917, M16248, K02121, X01087), papillomavirus (for example GenBank Accession No. NC\_001353), Herpes Simplex Virus (HSV for example GenBank Accession No. NC\_001345), and other viruses such as HTLV (for example GenBank Accession No. AJ430458). Due to the high sequence variability of many viral genomes, selection of siRNA molecules for broad therapeutic applications would likely involve the conserved regions of the viral genome. Nonlimiting examples of conserved regions of the viral genomes include but are not limited to 5'-Non Coding Regions (NCR), 3'- Non Coding Regions (NCR) and/or internal ribosome entry sites (IRES). siRNA molecules designed against conserved regions of various viral genomes will enable efficient inhibition of viral replication in diverse patient populations and may ensure the effectiveness of the siRNA molecules against viral quasi species which evolve due to mutations in the non-conserved regions of the viral genome.

Non-limiting examples of human genes that can be targeted using siRNA molecules of the invention using methods described herein include any human RNA sequence, for example those commonly referred to by Genbank Accession Number. These RNA sequences can be used to design siRNA molecules that inhibit gene expression and therefore abrogate diseases, conditions, or infections associated with expression of those genes. Such non-limiting examples of human genes that can be targeted using siRNA molecules of the invention include VEGFr (VEGFr-1 for example GenBank Accession No. XM\_067723, VEGFr-2 for example GenBank Accession No. AF063658), HER1, HER2, HER3, and HER4 (for example Genbank Accession Nos: NM\_005228, NM\_004448, NM\_001982, and NM\_005235 respectively), telomerase (TERT, for example GenBank Accession No. NM\_003219), telomerase RNA (for example GenBank Accession No. U86046), NFkappaB, Rel-A (for example GenBank Accession No. NM\_005228), NOGO (for example GenBank Accession No. AB020693), NOGOr (for example GenBank Accession No. XM\_015620), RAS (for example GenBank Accession No. NM\_004283), RAF (for example GenBank Accession No. XM\_033884), CD20 (for example GenBank Accession No. X07203), METAP2 (for example GenBank Accession No. NM\_003219), CLCA1 (for example GenBank Accession No. NM\_001285), phospholamban (for example GenBank Accession No. NM\_002667), PTP1B (for example GenBank Accession No. M31724), and others, for example, those shown in Table III.

5

10

15

20

25

The siNA molecule of the invention can also be used in a variety of agricultural applications involving modulation of endogenous or exogenous gene expression in plants using siNA, including use as insecticidal, antiviral and anti-fungal agents or modulate plant traits such as oil and starch profiles and stress resistance.

## 5 Example 24: Diagnostic uses

10

15

20

25

30

The siNA molecules of the invention can be used in a variety of diagnostic applications, such as in the identification of molecular targets (e.g., RNA) in a variety of applications, for example, in clinical, industrial, environmental, agricultural and/or research settings. Such diagnostic use of siNA molecules involves utilizing reconstituted RNAi systems, for example, using cellular lysates or partially purified cellular lysates. siNA molecules of this invention can be used as diagnostic tools to examine genetic drift and mutations within diseased cells or to detect the presence of endogenous or exogenous, for example viral, RNA in a cell. The close relationship between siNA activity and the structure of the target RNA allows the detection of mutations in any region of the molecule, which alters the base-pairing and three-dimensional structure of the target RNA. By using multiple siNA molecules described in this invention, one can map nucleotide changes, which are important to RNA structure and function in vitro, as well as in cells and tissues. Cleavage of target RNAs with siNA molecules can be used to inhibit gene expression and define the role of specified gene products in the progression of disease or infection. In this manner, other genetic targets can be defined as important mediators of the disease. These experiments will lead to better treatment of the disease progression by affording the possibility of combination therapies (e.g., multiple siNA molecules targeted to different genes, siNA molecules coupled with known small molecule inhibitors, or intermittent treatment with combinations siNA molecules and/or other chemical or biological molecules). Other in vitro uses of siNA molecules of this invention are well known in the art, and include detection of the presence of mRNAs associated with a disease, infection, or related condition. Such RNA is detected by determining the presence of a cleavage product after treatment with a siNA using standard methodologies, for example, fluorescence resonance emission transfer (FRET).

٠.,

In a specific example, siNA molecules that cleave only wild-type or mutant forms of the target RNA are used for the assay. The first siNA molecules (i.e., those that cleave

only wild-type forms of target RNA) are used to identify wild-type RNA present in the sample and the second siNA molecules (i.e., those that cleave only mutant forms of target RNA) are used to identify mutant RNA in the sample. As reaction controls, synthetic substrates of both wild-type and mutant RNA are cleaved by both siNA molecules to demonstrate the relative siNA efficiencies in the reactions and the absence of cleavage of the "non-targeted" RNA species. The cleavage products from the synthetic substrates also serve to generate size markers for the analysis of wild-type and mutant RNAs in the sample population. Thus, each analysis requires two siNA molecules, two substrates and one unknown sample, which is combined into six reactions. The presence of cleavage products is determined using an RNase protection assay so that full-length and cleavage fragments of each RNA can be analyzed in one lane of a polyacrylamide gel. It is not absolutely required to quantify the results to gain insight into the expression of mutant RNAs and putative risk of the desired phenotypic changes in target cells. The expression of mRNA whose protein product is implicated in the development of the phenotype (i.e., disease related or infection related) is adequate to establish risk. If probes of comparable specific activity are used for both transcripts, then a qualitative comparison of RNA levels is adequate and decreases the cost of the initial diagnosis. Higher mutant form to wildtype ratios are correlated with higher risk whether RNA levels are compared qualitatively or quantitatively.

All patents and publications mentioned in the specification are indicative of the levels of skill of those skilled in the art to which the invention pertains. All references cited in this disclosure are incorporated by reference to the same extent as if each reference had been incorporated by reference in its entirety individually.

One skilled in the art would readily appreciate that the present invention is well adapted to carry out the objects and obtain the ends and advantages mentioned, as well as those inherent therein. The methods and compositions described herein as presently representative of preferred embodiments are exemplary and are not intended as limitations on the scope of the invention. Changes therein and other uses will occur to those skilled in the art, which are encompassed within the spirit of the invention, are defined by the scope of the claims.

5

10

15

20

25

It will be readily apparent to one skilled in the art that varying substitutions and modifications can be made to the invention disclosed herein without departing from the scope and spirit of the invention. Thus, such additional embodiments are within the scope of the present invention and the following claims. The present invention teaches one skilled in the art to test various combinations and/or substitutions of chemical modifications described herein toward generating nucleic acid constructs with improved activity for mediating RNAi activity. Such improved activity can comprise improved stability, improved bioavailability, and/or improved activation of cellular responses mediating RNAi. Therefore, the specific embodiments described herein are not limiting and one skilled in the art can readily appreciate that specific combinations of the modifications described herein can be tested without undue experimentation toward identifying siNA molecules with improved RNAi activity.

The invention illustratively described herein suitably can be practiced in the absence of any element or elements, limitation or limitations that are not specifically disclosed herein. Thus, for example, in each instance herein any of the terms "comprising", "consisting essentially of", and "consisting of" may be replaced with either of the other two terms. The terms and expressions which have been employed are used as terms of description and not of limitation, and there is no intention that in the use of such terms and expressions of excluding any equivalents of the features shown and described or portions thereof, but it is recognized that various modifications are possible within the scope of the invention claimed. Thus, it should be understood that although the present invention has been specifically disclosed by preferred embodiments, optional features, modification and variation of the concepts herein disclosed may be resorted to by those skilled in the art, and that such modifications and variations are considered to be within the scope of this invention as defined by the description and the appended claims.

In addition, where features or aspects of the invention are described in terms of Markush groups or other grouping of alternatives, those skilled in the art will recognize that the invention is also thereby described in terms of any individual member or subgroup of members of the Markush group or other group.

5

10

15

20

# Table I

Targe	ge	Target Sequence	Seq	strand	RP#	Aliases		#
t Pos		CALILICGLICGUGGAAAUUCAACCU	₽ -	sense	30937	30937 ABCB1:120U21 siRNA	B unccuccuGGAAAuucAAcTT B	186
- 6		INCCUCIOCAUGAUGCUGGUGUUU	2	sense	30938	30938 ABCB1:620U21 siRNA	B ccucucAuGAuGcuGGuGuTT B	187
ABCB1 1867		CACGAUAGCUGAAAACAUUCGCU	<sub>6</sub>	sense	30939	stab04 ABCB1:1869U21 siRNA	B cGAuAGcuGAAAAcAuucGTT B	188
		AAAAIIGCAGCUGAUGAAUCCAAA	4	sense	30940	stab04 ABCB1:2336U21 siRNA	B AAuGcAGcuGAuGAAuccATT B	189
ABCB1 118		CAUUCCUCCUGGAAAUUCAACCU	-	antisense	30941	stab04 30941 ABCB1:138L21 siRNA (120C) stab05	GuuGAAuuuccAGGAGGAATsT	190
ABCB1 618		UUCCUCUCAUGAUGCUGGUGUUU	2	antisense	30942		AcAccAGcAucAuGAGAGGISI	<u>.</u>
ABCB1 1867		CACGAUAGCUGAAAACAUUCGCU	3	antisense	30943		cGAAuGuuucAGcuAucGTsT	192
		AAAAUGCAGCUGAUGAAUCCAAA	4	antisense		30944 ABCB1:2354L21 siRNA	uGGAuucAucAGcuGcAuuTsT	193
			-	gondo	31013	ABCB1:120U21 siRNA	UUCCUCCUGGAAAUUCAACTT	194
-+	-+	CAUUCCUCCUGGAAAUUCAACCU	- 2	sense	31014	31014 ABCB1:620U21 siRNA	ccucucaugaugeuguii	3 8
	-+-	CACCALIACTIGAAAACAUUCGCU	3	sense	31015	31015 ABCB1:1869U21 siRNA	CGAUAGCUGAAAACAOCCGTT	197
		AAAAIIGCAGCIIGAIIGAAUCCAAA	4	sense	31016	ABCB1:2336U21 siRNA	AAUGCAGCUGAUGAACCATT	138
ABCB1 23,	118 C	CAUUCCUCCUGGAAAUUCAACCU	1	antisense		31089 ABCB1:138L21 siRNA	GUUGAAOOOCCAGGAGGAGG	
	-	III CELICITO POR ANGENGO POR PORTO P	2	antisense		31090 ABCB1:638L21 siRNA	ACACCAGCAUCAUGAGAGGTT	199
	— <del> </del>	CACGALIAGCIIGAAAACAUUCGCU	3	antisense		(620C) 31091 ABCB1:1887L21 siRNA	CGAAUGUUUCAGCUAUCGTT	200
		AAAAIIGCAGCIJGAUGAAUCCAAA	4	antisense	31092		UGGAUUCAUCAGCUGCAUUTT	201
			LC.	sense	30721	(2336C) 30721 ADORA1:921U21 siRNA	B uucGAGAAGGucAucAGcATT B	202
ADORA 91	919 	AGUUCGAGAAGGUCAUCAGCAGG	,	3		stab04	B ccAGGuGucuAGAGGCAAcTT B	203
ADORA 1621		GACCAGGUGUCUAGAGGCAACAG	9	sense	30722	Stab04	B CARCOCCOLINA ACCIONATOR A TITE	204
ADORA 18	1819 G	GGACCAAGCUUAAGGAGGAGA	7	seuse	3072:	30723 ADORA1:1821U21 siRNA stab04	B Accadeculaaccandadouring	, R
ADORA 27	2773 G	GUCGGUUGACCUUCUGAACAUGA	80	seuse	† _	30724 ADORA1:2775U21 siRNA etah04	B cGGuuGAccuucucAAcAu I I B	3 8
			L	anticane	3072	20725 ADORAT:939L21 siRNA	uGcuGAuGAccuncucGAATsT	202

,								
- 000			-			(921C) stab05		
ADORA 1			9	antisense 30726	30726	ADORA1:1641L21 siRNA (1623C) stab05	GuuGccucuAGAcAccuGGTsT	202
ADORA 1	1819		7	antisense	30727		uccucuuAAGcuuGGuTsT	208
ADORA 1	2773	GUCGGUUGACCUUCUGAACAUGA	8	antisense		30728 ADORA1:2793L21 siRNA (2775C) stah05	AuGuucAGAAGGucAAccGTsT	209
ADORA 1	919	AGUUCGAGAAGGUCAUCAGCAUG	2	sense	31041	31041 ADORA1:921U21 SIRNA	UUCGAGAAGGUCAUCAGCATT	210
ADORA 1	1621	GACCAGGUGUCUAGAGGCAACAG	9	sense	31042	31042 ADORA1:1623U21 siRNA	CCAGGUGUCUAGAGGCAACTT	211
ADORA 1	1819	GGACCAAGCUUAAGGAGAGGAGA	7	sense	31043	31043 ADORA1:1821U21 siRNA	ACCAAGCUUAAGGAGAGGATT	212
ADORA 1	2773	GUCGGUUGACCUUCUGAACAUGA	ω	seuse	31044	31044 ADORA1:2775U21 siRNA	CGGUUGACCUUCUGAACAUTT	213
ADORA 1	919	AGUUCGAGAAGGUCAUCAGCAUG	co	antisense		31117 ADORA1:939L21 siRNA (921C)	UGCUGAUGACCUUCUCGAATT	214
ADORA 1			9	antisense		31118 ADORA1:1641L21 siRNA (1623C)	GUUGCCUCUAGACACCUGGTT	215
ADORA 1	1819	GGACCAAGCUUAAGGAGAGAGA	7	antisense	31119	ADORA1:1839L21 siRNA (1821C)	UCCUCUCCUUAAGCUUGGUTT	216
ADORA 1	2773		8	antisense		31120 ADORA1:2793L21 siRNA (2775C)	AUGUUCAGAAGGUCAACCGTT	217
b2a2	283	UGACCAUCAAUAAGGAAGAAGCC	9	sense	31594	31594 b2a2:283U21 siRNA	ACCAUCAAUAAGGAAGAGTT	218
b2a2	786	CCAUCAAUAAGGAAGAGCCCUU	10	sense	31595	31595 b2a2:286U21 siRNA	AUCAAUAAGGAAGAAGCCCTT	219
b2a2	282	CUGACCAUCAAUAAGGAAGAAGC	11	sense	31596	31596 b2a2:282U21 siRNA	GACCAUCAAUAAGGAAGAATT	220
b2a2	230	CAAUAAGGAAGAAGCCCUUCAGC	12	seuse	31597	31597 b2a2:290U21 siRNA	AUAAGGAAGAAGCCCUUCATT	221
b2a2	301	UGACCAUCAAUAAGGAAGAGCC	6	antisense	31598	31598  b2a2:301L21 siRNA (283C)	CUUCUUCCUUAUUGAUGGUTT	222
b2a2	304	CCAUCAAUAAGGAAGAAGCCCUU	10	antisense	31599	31599 b2a2:304L21 siRNA (286C)	GGGCUUCUUCCUUAUUGAUTT	223
b2a2	300	CUGACCAUCAAUAAGGAAGAGC	11	antisense	31600	31600 b2a2:300L21 siRNA (282C)	UNCUNCCUNAUNGANGGUCTT	224
b2a2	308	CAAUAAGGAAGAAGCCCUUCAGC	12	antisense	31601	31601 b2a2:308L21 siRNA (290C)	UGAAGGGCUUCUUCCUUAUTT	225
b3a2	356	UGGAUUUAAGCAGAGUUCAAAAG	13	sense	31602	31602 b3a2:356U21 siRNA	GAUUUAAGCAGAGUUCAAATT	226
b3a2	365	GCAGAGUUCAAAAGCCCUUCAGC	14	sense	31603	31603 b3a2:365U21 siRNA	AGAGUUCAAAAGCCCUUCATT	227
b3a2	364	AGCAGAGUUCAAAAGCCCUUCAG	15	sense	31604	31604 b3a2:364U21 siRNA	CAGAGUUCAAAAGCCCUUCTT	228
b3a2	357	GGAUUUAAGCAGAGUUCAAAAGC	16	sense	31605	31605 b3a2:357U21 siRNA	AUUUAAGCAGAGUUCAAAATT	229
b3a2	374	UGGAUUUAAGCAGAGUUCAAAAG	5	antisense	31606	31606 b3a2:374L21 siRNA (356C)	UUUGAACUCUGCUUAAAUCTT	230

231	232	233	234	235	236	237	238	239	240	241	242	243	244	245		246	247	248	249	250	251	252	253
UGAAGGGCUUUUGAACUCUTT	GAAGGGCUUUUGAACUCUGTT	UUUUGAACUCUGCUUAAAUTT	B uGGGuGAGGuuAccAAccATT B	B AccuuGGAcAuGGAAGAcuTT B	B uGGGAccuGcuAAGuGuGGTT B	uGGuuGGuAAccucAcccATsT	AGucuuccAuGuccAAGGuTsT	ccAcAcuuAGcAGGucccATsT	UGGGUGAGGUUACCAACCATT	ACCUUGGACAUGGAAGACUTT	UAACAUUGGUGCAAAGAUUTT	UGGGACCUGCUAAGUGUGGTT	UGGUUGGUAACCUCACCATT	AGUCUUCCAUGUCCAAGGUTT		AAUCUUUGCACCAAUGUUATT	CCACACUUAGCAGGUCCCATT	B uAAcAuuGGuGcAAAGAuuTT B	AAucuuuGcAccAAuGuuATsT	B uAAcAuuGGuGcAAAGAuuTT B	AAucuuuGcAccAAuGuuATsT	B uuAGAAAcGuGGuuAcAAuTT B	AuuGuAAccAcGuuucuAATsT
antisense 31607 b3a2:383L21 siRNA	31608 b3a2:382L21 siRNA	31609 b3a2:375L21 siRNA (357C)	30729 BACE:1492U21 siRNA stab04	30730 BACE:1755U21 siRNA stab04	30732 BACE:3585U21 siRNA stab04	30733 BACE:1510L21 siRNA (1492C) stab05			31005 BACE:1492U21 siRNA	31006 BACE:1755U21 siRNA	31007 BACE:2459U21 siRNA	31008 BACE:3585U21 siRNA		(1492C) RACE-17731 21 siRNA	(1755C)			31378 BACE:2459U21 siRNA stab04		31384 BACE:2459U21 siRNA	BACE:2477L21 siRNA	31390 BACE:2459U21 siRNA inv	31393 BACE:2477L21 siRNA (2459C) inv stab05
31607	31608	31609	30729	30730	30732	30733	30734	30736	31005	31006	31007	31008	31081	31082		31083	31084	31378	31381	31384	31387	31390	
antisense	antisense	antisense	seuse	seuse	sense	antisense	antisense	antisense	sense	sense	sense	sense	antisense	anticence	9611361130	antisense	antisense	seuse	antisense	seuse	antisense	seuse	antisense
14	15	16	17	18	19	17	8	19	17	18	8	19	17	4	2	20	19	20	20	20	20	20	20
GCAGAGUUCAAAAGCCCUUCAGC	AGCAGAGUUCAAAAGCCCUUCAG	GGAUUUAAGCAGAGUUCAAAAGC	AAUGGGUGAGGUUACCAACCAGU	UCACCUUGGACAUGGAAGACUGU	UAUGGGACCUGCUAAGUGUGGAA	AAUGGGUGAGGUUACCAACCAGU	UCACCUUGGACAUGGAAGACUGU	UAUGGGACCUGCUAAGUGUGGAA	AAHGGGHGAGGHIACCAACCAGU		CCUAACAUUGGUGCAAAGAUUGC	3583 HALIGGGACCUGCUAAGUGUGGAA	AAUGGGUGAGGUUACCAACCAGU		UCACCOOGGACACAO	CCUAACAUUGGUGCAAAGAUUGC	UAUGGGACCUGCUAAGUGUGGAA	CCUAACAUUGGUGCAAAGAUUGC	CCUAACAUUGGUGCAAAGAUUGC	CCUAACAUUGGUGCAAAGAUUGC	CCUAACAUUGGUGCAAAGAUUGC	CCUAACAUUGGUGCAAAGAUUGC	CCUAACAUUGGUGCAAAGAUUGC
383	382	375	1490	1753	3583	1490	1753	3583	1100		2457	- 1		1750	1/33	2457	3583	2457	2457	2457	2457	2457	2457
b3a2	b3a2	b3a2	BACE	BACE	BACE	BACE	BACE	BACE	BACE	RACE	BACE	RACE	BACE	L	BACE	BACE	BACE	BACE	BACE	BACE	BACE	BACE	BACE

1	
31565 CDK2:344U21 SIKNA	
seuse	
29	
ugn	

BCI 2	3220	3220 CAGGGAUGAUCAACAGGGUAGUG	24	antisense	31375	antisense 31375 BCL2:3240L21 siRNA	cccuAcuAGuuGucccAucTsT	717
			į		)	(3222C) inv stab11	B 11G11AG11GGGGUUCUAGGCATT B	278
CCND1	1628	GCUGUAGUGGGGUUCUAGGCAUC	23	seuse	30740	stab04		0.10
CCND1	2617	ACACACAAACCUUCUGCCUUUGA	56	seuse	30747	CCND1:2617U21 siRNA	B AcAcAAccuucuGccuuu I B	6/7
CCND1	3124	UCACAUUGUUUGCUGCUAUUGGA	27	sense	30748	30748 CCND1:3124U21 siRNA	B AcAuuGuuuGcuGcuAuuGTT B	280
CCND1	1646	GCUGUAGUGGGGUUCUAGGCAUC	25	antisense	30750	CCND1:1646L21 siRNA (1628C) stab05	uGccuAGAAcccAcuAcATsT	281
CCND1	2635	ACACACAAACCUUCUGCCUUUGA	26	antisense	30751	CCND1:2635L21 siRNA	AAAGGcAGAAGGuuuGuGuTsT	282
CCND1	3142	UCACAUUGUUUGCUGCUAUUGGA	27	antisense	30752		CAAUAGCAGCAAACAAUGuTsT	283
			96	gones	31009	31009 CCND1:695U21 siRNA	ACACUUCCUCUCCAAAAUGTT	284
SCND1	695	GAACACUUCCUCUCCAAAAUGCC	27,	Spinos	31010	31010 CCND1:1628U21 siRNA	UGUAGUGGGGUUCUAGGCATT	285
CCNC	1628	GCUGUAGGGGGGUCCAGGCAGGCAGGCAGGCAGGCAGGCA	3/8	sense	31011	31011 CCND1:2617U21 siRNA	ACACAAACCUUCUGCCUUUTT	286
	- 1	ACACACACACACACACACACACACACACACACACACAC	27	Sense	31012	CCND1:3124U21 siRNA	ACAUUGUUUGCUGCUAUUGT1	787
CCND1	713	GAACACUUCCUCUCCAAAAUGCC	78	antisense	31085	31085 CCND1:713L21 siRNA	CAUUUUGGAGAGGAAGUGUTT	288
CCND1	1646	GCUGUAGUGGGGUUCUAGGCAUC	25	antisense		31086 CCND1:1646L21 siRNA	UGCCUAGAACCCCACUACATT	289
CCND1	2635	ACACACAAACCUUCUGCCUUUGA	56	antisense	31087	31087 CCND1:2635L21 siRNA	AAAGGCAGAAGGUUUGUGUTT	290
CCND	3142	UCACAUUGUUUGCUGCUAUUGGA	27	antisense	31088	31088 CCND1:3142L21 siRNA	CAAUAGCAGCAAACAAUGUTT	291
					70070	(3124C)	R AcAcunccuccocAAAAuGTT B	292
CCND1	969	GAACACUUCCUCUCCAAAAUGCC	28	seuse	31304	Stab04	d Had week	COC
CCND1	695	GAACACUUCCUCCAAAAUGCC	78	seuse	31304	31304 CCND1:695U21 siRNA stab04	B AcAcunccucccAAAuG11 B	767
CCND1	695	GAACACUUCCUCUCCAAAAUGCC	78	seuse	31304	CCND1:695U21 siRNA stab04	B AcAcuuccucuccAAAAuGTT B	292
CCND1	713	GAACACUUCCUCUCCAAAAUGCC	28	antisense	31305	CCND1:713L21 siRNA (695C) stab()5	cAuuuuGGAGAGGAAGuGuTsT	293
CCND1	713	GAACACUUCCUCUCCAAAAUGCC	78	antisense	31305		cAuuuuGGAGAGGAAGuGuTsT	293
CCND1	695	GAACACUUCCUCUCCAAAAUGCC	28	seuse	31316		B GuAAAAccucuccuucAcATT B	294
CCND1	713	GAACACUUCCUCUCCAAAAUGCC	78	antisense	31317		uGuGAAGGAGAGGuuuuAcTsT	295
	-+		g	Sense	31565	31565 CDK2:344U21 siRNA	GGACACUGAGACUGAGGGUTT	296
CDK2	344	CUGGACACUGAGACUG	3	3				

297	298	299	300	301	302	303	304	305	306	307	308	309	310	311	312	313	314	315	316	317	318	319	320
22	53	53	Ж.	3(	3(	3(	3(	3	<u> </u>	3	9	3	3	3	က်	<u>ن</u>	ė i	က	'n	3.	3.	3	3,
AUCAAGCUAGCAGACUUUGTT	CUCACCUUCUAGUCUUGGCTT	ACGUUAGAUUUGCCGUACCTT	ACCCUCAGUCUCAGUGUCCTT	CAAAGUCUGCUAGCUUGAUTT	GCCAAGACUAGAAGGUGAGTT	GGUACGGCAAAUCUAACGUTT	B uGGucAcAGGAGAGAAGGcTT B	B AGAAGuuGGGcuAucAAuGTT B	B uucAGGGGAcAuGAGuuuuTT B	GccuucucucauGuGAccATsT	cAuuGAuAGcccAAcuucuTsT	AAAAcucAuGucccuGAATsT	UGGUCACAGGAGAGAGGCTT	AGAAGUUGGGCUAUCAAUGTT	AGGGUGAUGGAUUGGAGUUTT	UUCAGGGGACAUGAGUUUUTT	GCCUUCUCCUGUGACCATT	CAUUGAUAGCCCAACUUCUTT	AACUCCAUCCAUCACCCUTT	AAAACUCAUGUCCCCUGAATT	B AGGGuGAuGGAuuGGAGuuTT B	AAcuccAAuccAucAccuTsT	B uuGAGGuuAGGuAGuGGGATT B
31566 CDK2:654U21 siRNA	31567 CDK2:1245U21 siRNA	31568 CDK2:1428U21 siRNA	31569 CDK2:362L21 siRNA (344C)	31570 CDK2:672L21 siRNA (654C)	31571 CDK2:1263L21 siRNA (1245C)	31572 CDK2:1446L21 siRNA (1428C)	30753 CHEK1:371U21 siRNA stab04	30754 CHEK1:1351U21 siRNA stab04	30756 CHEK1:1880U21 siRNA stab04	30757 CHEK1:389L21 siRNA (371C) stab05	30758 CHEK1:1369L21 siRNA (1351C) stab05	30760 CHEK1:1898L21 siRNA (1880C) stab05	31001 CHEK1:371U21 SIRNA	31002 CHEK1:1351U21 siRNA	31003 CHEK1:1492U21 SIRNA	31004 CHEK1:1880U21 siRNA	31077 CHEK1:389L21 siRNA (371C)	31078 CHEK1:1369L21 siRNA (1351C)	31079 CHEK1:1510L21 siRNA (1492C)	31080 CHEK1:1898L21 siRNA (1880C)	31302 CHEK1:1492U21 siRNA stab04	31303 CHEK1:1510L21 siRNA (1492C) stab05	31314 CHEK1:1492U21 siRNA inv stab04
seuse	sense	sense	antisense	antisense	antisense	antisense	sense	seuse	sense	antisense	antisense	antisense	seuse	sense	sense	seuse	antisense	antisense	antisense	antisense	sense	antisense	sense
30	31	32	29	30	31	32	33	34	35	33	34	35	33	34	36	35	33	34	36	35	36	36	36
CCAUCAAGCUAGCAGACUUUGGA	cacucaccuucuagucuu	ACACGUUAGAUUUGCCGUACCAA	CUGGACACUGAGACUGAGGGUGU	CCAUCAAGCUAGCAGACUUUGGA	CACUCACCUUCUAGUCUUGGCCA	ACACGUUAGAUUUGCCGUACCAA	UAUGGUCACAGGAGAGGCAA	UGAGAAGUUGGGCUAUCAAUGGA	GUUUCAGGGGACAUGAGUUUUCC	UAUGGUCACAGGAGAGGCAA	UGAGAAGUUGGGCUAUCAAUGGA	GUUUCAGGGGACAUGAGUUUUCC	UAUGGUCACAGGAGAGGCAA	UGAGAAGUUGGGCUAUCAAUGGA	1490 UAAGGGUGAUGGAUUGGAGUUCA	GUUUCAGGGGACAUGAGUUUUCC	UAUGGUCACAGGAGAAGGCAA	UGAGAAGUUGGGCUAUCAAUGGA	UAAGGGUGAUGGAUUGGAGUUCA	GUUUCAGGGGACAUGAGUUUUCC	UAAGGGUGAUGGAUUGGAGUUCA	UAAGGGUGAUGGAUUGGAGUUCA	UAAGGGUGAUGGAUUGGAGUUCA
654	1245	1428	362	672	1263	1446	369	1349	1878	369	1349	1878	369	1349		1878	369	1349	1490	1878	1490	1490	1490
CDK2	CDK2	CDK2	CDK2	CDK2	CDK2	CDK2	CHEK1	CHEK1	CHEK1	CHEK1	CHEK1	CHEK1	CHEK1	CHEK1	CHEK1	CHEK1	CHEK1	CHEK1	CHEK1	CHEK1	CHEK1	CHEK1	CHEK1

321	322	323	324	325	326	327	328	328	330	331	332	333	334	335	336
ucccAcuAccuAAccucAATsT	B UAACCUCGUACUGGUGCCUCC B	B GGAGGCACCAGUACGAGGUUA B	B AAACUCCAAGAUCCCCAAUCA B	B UGAUUGGGGAUCUUGGAGUUU B	B GCAAAAACCCUGUGAUUUCCU B	B AGGAAAUCACAGGUUUUUGC B	B UUGGUCAGUUUCUGGCAGUUC B	B GAACUGCCAGAAACUGACCAA B	B CCUCCGUGGUCAUGCUCCAAU B	B AUUGGAGCAUGACCACGGAGG B	UAACCUCGUACUGGUGCCUCCUU	GGAGGCACCAGUACGAGGUUAUU	AAACUCCAAGAUCCCCAAUCAUU	UGAUUGGGGAUCUUGGAGUUUUU	GCAAAACCCUGUGAUUUCCUUU
CHEK1:1510L21 siRNA	RPI 21550 EGFR 3830L23 AS as siRNA Str 1 (sense)	antisense 25228 RPI 21550 EGFR 3830L23 AS as siRNA Str 2 (antisense)	RPI 21549 EGFR as siRNA Str 2 (antisense)		RPI 21545 EGFR as siRNA Str 2 (antisense)	25234 RPI 21545 EGFR as siRNA Str 1 (sense)	25235 RPI 21543 EGFR as siRNA Str 2 (antisense)		25249 RPI 21550 EGFR 3830L23 AS as siRNA Str 4 Concel Inverted Control	25250 RPI 21550 EGFR 3830L23 AS as siRNA Str 1 (sense) Inverted Control		1 (Sense) +20 overnang 25805 RPI 21550 EGFR 3830L23 AS as siRNA Str 2 (antisense) +2U	antisense 25806 RPI 21549 EGFR as siRNA Str 2 (antisense)+	25807 RPI 21549 EGFR 3 as siRNA Str 1 (sense)+2U	25810 RPI 21545 EGFR as siRNA Str 2
31315	25227	25228	25229	25230	25233	25234	25235	25236	25249	25250	25804		25806	25807	2581(
antisense 31315	sense	antisense	antisense	sense	antisense	seuse	antisense	sense	sense	sense	seuse	antisense	antisense	seuse	antisense
36	37	38	39	40	41	42	43	44	38	45	37	38	39	40	41
CHEK1 1490   UAAGGGUGAUGGAUUGGAGUUCA	UAACCUCGUACUGGUGCCU	Accucenacueeueccucc	AUUGGGGAUCUUGGAGUUU	UGAUUGGGGAUCUUGGAGU	GAAAUCACAGGGUUUUUGC	AGGAAAUCACAGGGUUUUU	ACUGCCAGAAACUGACCAA	GAACUGCCAGAAACUGACC	Accuceuacueeueccucc	AGGCACCAGUACGAGGUUA	UAACCUCGUACUGGUGCCU	ACCUCGUACUGGUGCCUCC	AUUGGGGAUCUUGGAGUUU	UGAUUGGGGAUCUUGGAGU	GAAAUCACAGGGUUUUUGC
1490	3828								3828	3828	3828				
CHEK1	EGFR	EGFR	EGFR	EGFR	EGFR	EGFR	EGFR	EGFR	EGFR	EGFR	EGFR	EGFR	EGFR	EGFR	EGFR

UUUGCUU 337	AGUUCUU 338													
AGGAAAUCACAGGGUUUUGCUU	UNGGUCAGUUCUGGCAGUUCUU		GAACUGCCAGAAACUGACCAAUU	GAACUGCCAGAAACUGACCAAUU B UAACCUCGUACUGGUGCCUCCUU B	GAACUGCCAGAAACUGACCAAUU  B UAACCUCGUACUGGUGCCUCCUU B  B GGAGGCACCAGUACGAGGUUAUU B	GAACUGCCAGAAACUGACCAAUU  B UAACCUCGUACUGGUGCCUCCUU E  B GGAGGCACCAGUACGAGGUUAUU E  B AAACUCCAAGAUCCCCAAUCAUU B	GAACUGCCAGAAACUGACCAAUU  B UAACCUCGUACUGGUGCCUCCUU B  B GAGGCACCAGUACGAGGUUAUU B  B AAACUCCAAGAUCCCCAAUCAUU B  B UGAUUGGGGAUCUUGGAGUUUUU B	GAACUGCCAGAAACUGACCAAUU  B UAACCUCGUACUGGUGCCUCCUU B  B GAAGUCCAAGUCCCCAAUCAUU B  B UGAUUGGGGAUCUUGGAGUUUUU B  B GCAAAAACCCUGUGAUUUCCUUU B	GAACUGCCAGAAACUGACCAAUU  B UAACCUCGUACUGGUGCCUCCUU B  B GGAGGCACCAGUACGAGGUUUUU B  B UGAUUGGGGAUCUUGGAGUUUUU B  B GCAAAAACCCUGUGAUUUCCUUU B  B AGGAAAUCACAGGUUUUUGCUU B	GAACUGCCAGAAACUGACCAAUU  B UAACCUCGUACUGGUGCCUCCUU B  B GGAGGCACCAGUACGAGGUUAUU B  B AAACUCCAAGAUCCCCAAUCAUU B  B GGAAAAACCCUGUGAUUUCCUUU B  B AGGAAAUCACAGGUUUUUGCUU B  B AGGAAAUCACAGGGUUUUUGCUU B	GAACUGCCAGAAACUGACCAAUU  B UAACCUCGUACUGGUGCCUCCUU I  B GGAGGCACCAGUACGAGGUUAUU B  B AAACUCCAAGAUCCCCAAUCAUU B  B GGAAAAACCCUGUGAUUUUCCUUU B  B AGGAAAUCACAGGGUUUUUGCUU B  B AGGAAAUCACAGGGUUUUUGCUU B  B GGAAAUCACAGGGUUUUUGCUU B	GAACUGCCAGAAACUGACCAAI  B UAACCUCGUACUGGUGCCUCCI  B GAGGGCACCAGUACCAAUCAU  B AAACUCCAAGAUCCCCAAUCAU  B GAAAAACCCUGUGAUUUCCUU  B AGGAAAUCACAGGGUUUUUGCU  B AGGAAAUCACAGGGUUUUUGCU  B AGGAAAUCACAGGGUUUUUGCU  B AGGAAAUCACAGGAAACUGACCAAU	GAACUGCCAGAAACUGACCAAL  B UAACCUCGUACUGGUGCCUCCU  B GAACUCCAAGAUCCCCAAUCAU  B UGAUUGGGGAUCUUGGAGUUUU  B GAAAAACCCCUGUGAUUUUGCU  B AGGAAAACCCAGAAACUGACCAAUCU  B AGGAAAUCACAGGAAACUGACCAAUI  B GAACUGCCAGAAACUGACCAAUI  B GAACUGCCAGAAACUGACCAAUI  B GAACUGCCAGAAACUGACCAAUI  B GAACUGCCAGAAACUGACCAAUI  B GAACUGCCAGAAACUGACCAAUI  B GAACUGCCAGAAACUGACCAAUI	GAACUGCCAGAAACUGACCAAA  B UAACCUCGUACUGGUGCCUCC  B GAACUCCAAGAUCCCCAAUCAU  B AAACUCCAAGAUCCCCAAUCAU  B GAAAAACCCCUGUGAUUUUCCUU  B AGGAAAUCACAGGGUUUUUGCU  B GAACUGCCAGAAACUGACCAAU  B GAACUGCCAGAAACUGACCAAU  B GAACUGCCAGAAACUGACCAAU  B GAACUGCCAGAAACUGACCAAU  B GAACUGCCAGAAACUGACCTT B  B GAACUGCCAGAAACUGACCTT B
	as		25	SU Str	as as e)+2U RNA Str erhang	as as e)+2U RNA Str arhang RNA Str as as ense)+	as as hand as RNA Str RNA Str RNA Str arbang as ense)+	as RNA Str RNA Str as ense)+ as as ense)+ as as ense)+ as ense)+ as enhang	as se)+2U RNA Str RNA Str RNA Str RNA Str as ense)+ as erhang erhang as sas erhang as sas so)+2U	as RNA Str RNA Str RNA Str RNA Str as ense)+ as as ense)+ as a	as se)+2U RNA Str and as as ense)+ as as erhang as as erhang as as se)+2U as as Erhang as	as as enhang enha	as as enhang enha	as as hang as enhang as en
	SIKNA Str 2	25813 RPI 21543 EGFR as siRNA Str 1 (sense)+2U	overhand	overhang 25824 RPI 21550 EGFR 3830L23 AS as si 1 (sense) +211 ove	overhang sense 25824 RPI 21550 EGFR 3830L23 AS as si 1 (sense) +2U ove antisense 25825 RPI 21550 EGFR 3830L23 AS as sil 2 (antisense) +2U	overhang 25824 RPI 21550 EGFR 3830L23 AS as siRN 1 (sense) +2U overh 25825 RPI 21550 EGFR 3830L23 AS as siRN 2 (antisense) +2U overhang 25826 RPI 21549 EGFR as siRNA Str 2 (antisense) 2U overhang	25824 RPI 21550 EGFR 3830L23 AS as siRNA 1 (sense) +2U overhar 25825 RPI 21550 EGFR 3830L23 AS as siRNA 2 (antisense) +2U 20verhang 25826 RPI 21549 EGFR as siRNA Str 2 (antisense 2U overhang 2U overhang 2U overhang 2U overhang 2U overhang 2U overhang	overhang 25824 RPI 21550 EGFR 3830L23 AS as siRN 1 (sense) +2U overha 25825 RPI 21550 EGFR 3830L23 AS as siRN 2 (antisense) +2U overhang 25826 RPI 21549 EGFR as siRNA Str 2 (antisense)+2U overhang 25827 RPI 21549 EGFR 3a siRNA Str 1 (sense)+ overhang 25830 RPI 21545 EGFR as siRNA Str 2 overhang 25830 RPI 21545 EGFR as siRNA Str 2 (antisense)+2U overh	THE WORLD STEEL ST	STATE OF STATE SOIL SOIL SOIL SOIL SOIL	THE WOLL WITH WALL WOLL WOLL WOLL WOLL	A STAN STAN STAN STAN STAN STAN SOLD SOLD STAN STAN STAN STAN STAN STAN STAN STAN	the man and a superior and a superio	St mist mist miss as a constant and
antisense 25812		sense 258	1-		sense 2582	antisense 2582 antisense 2582			sense 25825 sense 25826 sense 25827 ense 25837		sense 2582 nse 2582 ense 2583 ense 25833 sense 25833	sense 2582 sense 2582 nse 2583 ense 2583 ense 25833 se 25833	sense 2582 sense 2582 nse 2583 ense 2583 ense 2583 se 25833 se 30705 se 30706	sense 25826 sense 25826 ense 25837 ense 25833 se 30705 se 30705 se 30707
_	43 antise	44 sen	37 sense		38 antise									
0000   42				_							4	4 4	4 4 4	4 4 4
	ACUGCCAGAAACUGACCAA	GAACUGCCAGAAACUGACC	UAACCUCGUACUGGUGCCU		ACCUCGUACUGGUGCCUCC	ACCUCGUACUGGUGCCUCC	ACCUCGUACUGGUGCCUCC AUUGGGGAUCUUGGAGUU UGAUUGGGGAUCUUGGAGU	ACCUCGUACUGGUGCCUCC AUUGGGGAUCUUGGAGU UGAUUGGGGAUCUUGGAGU	ACCUCGUACUGGUGCCUCC  AUVGGGGGAUCUUGGAGU  UGAUUGGGGGAUCUUGGAGGU  GAAAUCACAGGGUUUUUGC	ACCUCGUACUGGUGCCUCC AUUGGGGGAUCUUGGAGUU UGAUUGGGGAUCUUGGAGUU GAAAUCACAGGGUUUUUU AGGAAAUCACAGGGUUUUUU AGGAAAUCACAGGGUUUUUU	ACCUCGUACUGGUGCCUCC AUUGGGGAUCUUGGAGUU UGAUUGGGGAUCUUGGAGUU GAAAUCACAGGGUUUUUU AGGAAAUCACAGGGUUUUUU AGGAAAUCACAGGGUUUUUU GAACUGCCAGAAACUGACC	ACCUCGUACUGGUGCCUCC AUUGGGGAUCUUGGAGUU UGAUUGGGGAUCUUGGAGUU GAAAUCACAGGGUUUUUUGC GAAAUCACAGGGUUUUUUU AGGAAAUCACAGGAAACUGACC GAACUGCCAGAAACUGACC	ACCUCGUACUGGUGCCUCC AUUGGGGAUCUUGGAGUU UGAUUGGGGAUCUUGGAGUU GAAAUCACAGGGUUUUUUGC GAAAUCACAGGGUUUUUU ACUGCCAGAAACUGACC GAACUGCCAGAAACUGACC AGGAAAUCACAGGGUUUUU	ACCUCGUACUGGUGCCUCC AUUGGGGAUCUUGGAGUU UGAUUGGGGAUCUUGGAGUU GAAAUCACAGGGUUUUUU AGGAAAUCACAGGGUUUUUU AGGAAAUCACAGAAACUGACC GAACUGCCAGAAACUGACC AGGAAAUCACAGGGUUUUU GUUCCGUGAGUUGAUCAUC
EGFR	<u></u>		3828	_							,		.	,
	EGFR	EGFR	EGFR	LC-LC-LC-LC-LC-LC-LC-LC-LC-LC-LC-LC-LC-L		EGFR	EGFR EGFR	EGFR EGFR EGFR	EGFR EGFR EGFR	EGFR EGFR EGFR EGFR	EGFR EGFR EGFR EGFR	EGFR EGFR EGFR EGFR EGFR	EGFR EGFR EGFR EGFR	

$\infty$	
4	
_	

352		353	354	355	356	357	358	359	360	26.1	100	362	263	202	351	255	CCS	364	365	3	366	7967	705	368	369	0.00	3/0	371
Talamayouth	Gencycanacia	AAAAcccuGuGAuuuccuTsT	GAuGAucAcucAcGGAAcTsT	uGGAGucuGuAGGAcuuGGTsT	TUDAGEAAACIIGACCTT	GAACUGCCAGAACUCACA	GIIICGGIGAGUUGAUCAUCTT	CCAAGUCCUACAGACUCCATT	GGUCAGUUCUGGCAGUUCTT	THE COLUMN THE PROPERTY OF THE	AAAAACCCUGUGAUUUCCUII	GAUGAUCAACUCACGGAACTT		UGGAGUCUGUAGGACUUGGTI	B ccAAGuccuAcAGAcuccATT B	H 1000	uGGAGucuGuAGGAcuuGG1s1	B AccucAGAcAuccuGAAccTT B	Lot.	GGuucAGGAuGucucAGGuisi	B G. GAALGGGUCAAGGAACUTT B		B GGAAcuGuGcAAGAuGAcc11 B	B AAGcuGcucAAccAucuccTT B	R cAAccAucuccuuccAcAGTT B		AGuuccuuGAGccAuucAc1s1	GGucAucunGcAcAGuuccTsT
stab04	9 EGFR:819L21 siRNA		=	(3066C) stabub 2 EGFR:3172L21 siRNA		30985 EGFR:801U21 siRNA	30986 EGFR:1382U21 SIKNA	30987 EGFR:30000Z1 SINIVA	30988 EGFK:31340Z1 SINIA	antisense 31001 EGFR.019E21 SING (801C)	31062 EGFR:1400L21 siRNA	(1382C)	(3066C)	31064 EGFR:3172L21 siRNA	3134C)	stab04		(3154C) Stabus 19 ECER-31541/21 siRNA inv	stab04	,	(3154C) Inv stabub	30761 ERG2:244UZ1 SIRIVA stab04	62 ERG2:519U21 siRNA	30763 ERG2:761U21 siRNA	stab04	30764 ERGZ:7690z 1 sirkin stab04	30765 ERG2:262L21 siRNA	antisense 30766 ERG2:537L21 siRNA
	30208	9 30710	e. 3071	e 3071	3			$\neg$	303	9 3100 -			se 31063			31300	se 31301	24249		antisense 31313			30762					se 307
	antisense 30709	antisense	antisense, 30711	antisense 30712		seuse	seuse	sense	seuse	antisens	antisense		antisense	antisense		sense	antisense		seuse	antisen		seuse	seuse	sense		sense	antisense	antiser
	44	42	46	47		44	42	46	1	44	42		46	47	į	47	47	į	74	47		48	49	20		51	48	46
	GAACUGCCAGAAACUGACC	AGGAAAUCACAGGGUUUUU	GIIICCGUGAGUUGAUCAUC	VJJIIJVJVJVIIGOTIGOTIGOTIGOTIGOTIGOTIGOTIGOTIGOTIGO	CCAAGUCCUACAGACOCCA	GAACUGCCAGAAACUGACC	AGGAAAUCACAGGGUUUUU	GUUCCGUGAGUUGAUCAUC	CCAAGUCCUACAGACUCCA	GAACUGCCAGAAACUGACC	AGGAAAUCACAGGGUUUUU		GUUCCGUGAGUUGAUCAUC	CCAAGUCCUACAGACUCCA	4	CCAAGUCCUACAGACUCCA	CCAAGUCCUACAGACUCCA		CCAAGUCCUACAGACUCCA	CCAAGUCCUACAGACUCCA		AGGUGAAUGGCUCAAGGAACUCU	AAGGAACUGUGCAAGAUGACCAA	CAAACCHGCHCAACCAHCHCCOUU		CUCAACCAUCUCCUUCCACAGUG	AGGI GAAUGGCUCAAGGAACUCU	AAGGAACUGUGCAAGAUGACCAA
-	799	1380	3064		3152	7007	38	3064	3152	799	1380	2	3064	3152		3152	3152	1	3152	3452	1	242	517		66 /	797	242	
	EGFR	-+-		-+	EGFR	FCFR	FGFR			EGFR	FOFF	- 5 1	EGFR	EGFR	;	EGFR	FCFR	- 5	EGFR	CCED	5	ERG2	FRG2		EKGZ	ERG2	2000	ERG2

372	373	374	375	376	377	378	379	380	381	382	383	384	385	386	387	388	386	390	391	392	393	394	395	396
GGAGAuGGuuGAGcAGcuuTsT	cuGuGGAAGGAGAuGGuuGTsT	GUGAAUGGCUCAAGGAACUTT	GGAACUGUGCAAGAUGACCTT	AAGCUGCUCAACCAUCUCCTT	CAACCAUCUCCUUCCACAGTT	AGUUCCUUGAGCCAUUCACTT	GGUCAUCUUGCACAGUUCCTT	GGAGAUGGUUGAGCAGCUUTT	CUGUGGAAGGAGGUGGUGTT	CAUGCGACUGAGACAGCUCTT	ACAUCCUGACUUCUGUGAGTT	GAUGAUGAUGGAGACGTT	ACAAUUUCUGUGCCAUUGCTT	GAGCUGUCACACACAUGTT	CUCACAGAAGUCAGGAUGUTT	CGUCUCCAUCAUCAUCTT	GCAAUGGCACAGAAAUUGUTT	CSUSGSASGSUUUAAAAGGCACCCTST	CSASASCSCSACAAAAUACAACAATST	CSCSUSGSGSAAAGAAUCAAAACCTST	GSCSASASGSGGGGCCUCUGAUGTST	GSGSGSUSGSCCUUUUAAACUCAGTST	USUSGSUSUSGUAUUUUGUGGUUGTST	GSGSUSUSUSUGAUUCUUUCCAGGTST
antisense 30767 ERG2:779L21 siRNA (761C) stab05	antisense 30768 ERG2:787L21 siRNA (769C) stab05	sense 31045 ERG2:244U21 siRNA	sense 31046 ERG2:519U21 siRNA	sense 31047 ERG2:761U21 siRNA	sense 31048 ERG2:769U21 siRNA	antisense 31121 ERG2:262L21 siRNA (244C)	antisense 31122 ERG2:537L21 siRNA (519C)	antisense 31123 ERG2:779L21 siRNA (761C)	antisense 31124 ERG2:787L21 siRNA (769C)	sense 31416 EZH2:203U21 siRNA	sense 31417 EZH2:340U21 siRNA	sense 31418 EZH2:690U21 siRNA	sense 31419 EZH2:1495U21 siRNA	antisense 31420 EZH2:221L21 siRNA (203C)	antisense 31421 EZH2:358L21 siRNA (340C)	antisense 31422 EZH2:708L21 siRNA (690C)	antisense 31423 EZH2:1513L21 siRNA (1495C)	sense 29694 FLT1:349U21 siRNA stab01	sense 29695 FLT1:2340U21 siRNA stab01	sense 29696 FLT1:3912U21 siRNA stab01	sense 29697 FLT1:2949U21 siRNA stab01	antisense 29698 FLT1:369L21 siRNA (349C) stab01	antisense 29699 FLT1:2358L21 siRNA (2340C) stab01	antisense 29700 FLT1:3932L21 siRNA
50 antis	51 antis	48 se	49 se	50 se	51 se	48 antis	49 antis	50 antis	51 antis	52 sel	53 sel	54 sel	55 sel	52 antis	53 antis	54 antis	55 antis	56 ser	57 ser	58 ser	59 ser	56 antis	57 antis	58 antis
GAAAGCUGCUCAACCAUCUCCUU	CUCAACCAUCUCCUUCCACAGUG	AGGUGAAUGGCUCAAGGAACUCU 4	AAGGAACUGUGCAAGAUGACCAA 4	GAAAGCUGCUCAACCAUCUCCUU	CUCAACCAUCUCCUUCCACAGUG	ļ	AAGGAACUGUGCAAGAUGACCAA 4	GAAAGCUGCUCAACCAUCUCCUU	CUCAACCAUCUCCUUCCACAGUG	UACAUGCGACUGAGACAGCUCAA 5	GCACAUCCUGACUUCUGUGAGCU	ACGAUGAUGAUGGAGACGAU	UGACAAUUUCUGUGCCAUUGCUA	UACAUGCGACUGAGACAGCUCAA	GCACAUCCUGACUUCUGUGAGCU	ACGAUGAUGAUGAUGGAGACGAU 5	UGACAAUUUCUGUGCCAUUGCUA	AACUGAGUUUAAAAGGCACCCAG 5	AACAACCACAAAAUACAACAAGA 5	AGCCUGGAAAGAAUCAAAACCUU	AAGCAAGGAGGCCUCUGAUGGU 5	AACUGAGUUUAAAAGGCACCCAG 5	AACAACCACAAAAUACAACAAGA 5	AGCCUGGAAAGAAUCAAAACCUU 5
759	767	242	517	759	797	242	517	759	792	201	338	889	1493	201	338	989	1493	347	2338	3910	2947	347	2338	3910
ERG2	ERG2	ERG2	ERG2	ERG2	ERG2	ERG2	ERG2	ERG2	ERG2	EZH2	EZH2	EZH2	EZH2	EZH2	EZH2	EZH2	EZH2	FLT1	FTT	FLT1	FLT1	FLT1	FLT1	FLT1

_	
⊂	>
v	ì
_	4

397	9	398	399	400	401	- I	402	403	404	405	406	407	P	408	409	410	144	- 3	412	5 2	414	415	416	417	418	
E TSTOBILIOUIDOGOVOVO CAROLINA CONTRACTOR CO	-		csAsAscscAcAAAAAAAAAAAAASTsT	GSCS118GSGAAAGAAucAAAAScscsTsT	-	- 1	GsGsGsUsGsCsCsUsUsUsAsAsAsCsUs CsAsGsTsT	UsUsGsUsGsG	SOSOSOSOS	SCSUSCSUS	USGSCSTST		UUGUUGUAUUUGUGGUAGAA	ASASCSASASCAUAAAACACCAACTST	GSUSUSGSGSUGUUUNUAUGUUGUUTST	TSTSUB QUANTO VAN A A A A A A A A A A A A A A A A A A	Asassasacauaaaacacasassas	GsUsUsGsGsUsGsUsBsUsAsUsGsUsUs SGsUsUsTsT	AGAACAUAAAACACCAAC	ບນຣຸບນຣຣບຣບປປປAປຣບນຣູນ	CAACCACAAAAUACAACAATI	UUGUUGUAUUUGUGGUUGLL	-		TJOHNGUIGHINING	
(3912C) stab01	701 FLT1:2969L21 siRNA (7949C) stab01	29702 FLT1:349U21 siRNA	29703 FLT1:2340U21 siRNA	stab03	29704 FLT1:3912UZ1 SIKINA stab03	949U21 siRNA	39L21 siRNA	siRNA	SIRNA	(3912C) stab02 29709 FLT1:2969L21 siRNA	(2949C) stab02	29981   FL   1:2340021 SIRNA Native	29982 FLT1:2358L21 siRNA	29983 FLT1:2342U21 siRNA	stab01 inv	29984 FELTI.2330E21 SINK	29985 FLT1:2342U21 siRNA stab03 inv	29986 FLT1:2358L21 siRNA	29987 FLT1:2340U21 siRNA inv	29988 FLT1:2358L21 siRNA	20075 El T1-2340121 SIRNA	30076 FLT1:2358L21 siRNA	(2340C)	30077 FLT1:2358L21 siRNA	(2340C) inv	antisense   30187   FLT1:2358L21 sIKNA
	antisense 29701	sense 29.	sense 29	_	sense 29	sense 29	antisense 29	antisense 29	antisense 29			sense   29	antisense 29	sense 29		antisense   2	sense 2	antisense 2	sense 2	antisense 2	十	sense 3		Sense		antisense
	59 a	56	57	<del>-  </del>	28	29	56	57 8	28	-		22	22	57	;	22	22	57	57	22	1	57		57	5	22
	AAGCAAGGAGGGCCUCUGAUGGU	AACUGAGUUUAAAAGGCACCCAG	A A COA A COA A A A A I I A C A A C A A A A A A A A	AACAACCACACACACACACACACACACACACACACACACA	AGCCUGGAAAGAAUCAAAACCUU	AAGCAAGGAGGCCUCUGAUGGU	AACUGAGUUUAAAAGGCACCCAG			AGCCOGGGGGGG	4	AACAACCACAAAAUACAACAAGA	AACAACCACAAAAUACAACAAGA			AACAACCACAAAAUACAACAAGA	AACAACCACAAAAUACAACAAGA	AACAACCACAAAAUACAACAAGA				AACAACCACAAAAUACAACAAGA			AACAACCACAAAAUACA	3 AACAACCACAAAAUACAACAAGA
	2947	347		2338	3910	2947	347	0000	2330	01.65	2947	2338	2338	9,00	2340	2338	2340	2338	_			++	7338	2340		2338
	FLT1	17.		F. 1.1	FLT1	FLT1	FI T1	F		FL11	FLT1	FLT1	FI T1			FLT1	FLT1	FLT1	FLTA	FLT1	i i	FLT1	FE3	FIT	FLT1	FLT1

RNA						L	0 11 2 10 (00/60)		
2338         AACAACCACAAAUACAACAAGA         57         antisense         30193         FLT1.2338L21 siRNA           2338         AACAACCACAAAUACAACAAGA         57         sense         30199         FLT1.2340U21 siRNA           2338         AACAACCACAAAUACAACAAGA         57         sense         30199         FLT1.2340U21 siRNA           2338         AACAACCACAAAUACAACAAGA         57         antisense         30349         FLT1.2358L21 siRNA           2338         AACAACCACAAAUACAACAAGA         57         antisense         30344         FLT1.2358L21 siRNA           2338         AACAACCACAAAUACAACAAGA         57         antisense         30346         FLT1.2358L21 siRNA           2338         AACAACCACAAAUACAACAAGA         57         antisense         30346         FLT1.2358L21 siRNA           2338         AACAACCACAAAUACAACAAGA         57         antisense         30346         FLT1.2358L21 siRNA     <	FLT1	<del> </del>		22	antisense	30190	(2340C) Z-F U,C FLT1:2358L21 siRNA	unGunGuAuunuGuGGuuGXX	419
1 2338         AACAACCACAAAUACAACAGA         57 sense         30196 [LT11.2340121 siRNA]           2338         AACAACCACAAAAUACAACAGA         57 sense         30196 [LT11.2340121 siRNA]           2338         AACAACCACAAAAUACAACAGA         57 antisense         30340 [LT1.2358121 siRNA]           2338         AACAACCACAAAAUACAACAGA         57 antisense         30340 [LT1.2358121 siRNA]           2338         AACAACCACAAAAUACAACAGA         57 antisense         30342 [LT1.2358121 siRNA]           2338         AACAACCACAAAAUACAACAAGA         57 antisense         30342 [LT1.2358121 siRNA]           2338         AACAACCACAAAUACAACAAGA         57 antisense         30342 [LT1.2358121 siRNA]           2338         AACAACCACAAAUACAACAAGA         57 antisense         30345 [LT1.2358121 siRNA]           2338         AACAACCACAAAUACAACAAGA         57 antisense         30346 [LT1.2358121 siRNA]           2338         AACAACCACAAAUACAACAAGA         57 antisense         30346 [LT1.2358121 siRNA]           2338         AACAACCACAAAUACAACAAGA         57 antisense         30346 [LT1.2358121 siRNA]           2338         AACAACCACAAAUACAACAAGA         57 antisense         30446 [LT1.2358121 siRNA]           2338         AACAACCACAAAUACAAAUACAACAAGA         57 antisense         307777 [LT1.11340121 siRNA]           4773	FLT1		l	22	antisense		FLT1:2358L21 siRNA	uuGuuGuAuuuuGuGGuuGZZ	420
2338         AACAACCACAAAUACAACAGA         57         sense         30199         FLT1.2340U21 siRNA           2338         AACAACCACAAAUACAACAGA         57         antisense         30349         FLT1.2358L2 siRNA           2338         AACAACCACAAAUACAACAGAGA         57         antisense         30341         FLT1.2358L2 siRNA           2338         AACAACCACAAAUACAACAGAGA         57         antisense         30341         FLT1.2358L2 siRNA           2338         AACAACCACAAAUACAACAGAGA         57         antisense         30343         FLT1.2358L2 siRNA           2338         AACAACCACAAAUACAACAGAGA         57         antisense         30343         FLT1.2358L2 siRNA           2338         AACAACCACAAAUACAACAGAGA         57         antisense         30345         FLT1.2358L2 siRNA           2338         AACAACCACAAAUACAACAGAGA         57         antisense         30346         FLT1.2358L2 siRNA           2338         AACAACCACAAAUACAACAAGAGA         57         antisense         30346         FLT1.2358L2 siRNA           2338         AACAACCACAAAUACAACAAGAGA         57         antisense         30346         FLT1.2358L2 siRNA           2338         AACAACCACAAAUACAAAUACAACAGAGA         57         antisense         30779         FLT1.1350L2 siRNA </td <td>FLT</td> <td><u> </u></td> <td>1</td> <td>22</td> <td>seuse</td> <td>30196</td> <td>FLT1:2340U21 siRNA</td> <td>B cAAccACAAAAUACAACAATT B</td> <td>421</td>	FLT	<u> </u>	1	22	seuse	30196	FLT1:2340U21 siRNA	B cAAccACAAAAUACAACAATT B	421
2338         AACAACCACAAAAUACAACAAGA         57         antisense         30340   ELT1.236B.21 siRNA           2338         AACAACCACAAAAUACAACAAGA         57         antisense         30341   ELT1.235B.21 siRNA           2338         AACAACCACAAAAUACAACAAGA         57         antisense         30342   ELT1.235B.21 siRNA           2338         AACAACCACAAAAUACAACAAGA         57         antisense         30342   ELT1.235B.21 siRNA           2338         AACAACCACAAAAUACAACAAGA         57         antisense         30348   ELT1.235B.21 siRNA           2338         AACAACCACAAAAUACAACAAGA         57         antisense         30346   ELT1.235B.21 siRNA           2338         AACAACCACAAAAUACAACAAGA         57         antisense         30346   ELT1.235B.21 siRNA           2338         AACAACCACAAAAUACAACAAGA         57         antisense         30346   ELT1.235B.21 siRNA           2338         AACAACCACAAAUACAACAAGA         57         antisense         3046   ELT1.235B.21 siRNA           2338         AACAACCACAAAUACAACAAGA         57         antisense         304777   ELT1.1144U21 siRNA           2338         AACAACCACAAAUACAACAAGA         60         sense         30777   ELT1.4753U21 siRNA           2338         AACAACCACAAAAUACAAAAGAAAAGAAAAGAAAGAAAGA	FLT		<u> </u>	22	seuse	30199	FLT1:2340U21 siRNA	CAACCACAAAAUACAACAATT	422
2338         AACAACCACAAAUACAACAAGA         57         antisense         30341   FLT1:2358L21 siRNA           2338         AACAACCACAAAUACAACAAGA         57         antisense         30342   FLT1:2358L21 siRNA           2338         AACAACCACAAAAUACAACAAGA         57         antisense         30344   FLT1:2358L21 siRNA           2338         AACAACCACAAAAUACAACAAGA         57         antisense         30344   FLT1:2358L21 siRNA           2338         AACAACCACAAAAUACAACAAGA         57         antisense         30344   FLT1:2358L21 siRNA           2338         AACAACCACAAAUACAACAAGA         57         antisense         30346   FLT1:2358L21 siRNA           2338         AACAACCACAAAUACAACAAGA         57         antisense         30346   FLT1:2358L21 siRNA           2338         AACAACCACAAAUACAACAAGA         57         antisense         30446   FLT1:2358L21 siRNA           2338         AACAACCACAAAUACAACAAGA         57         antisense         3046   FLT1:2358L21 siRNA           2338         AACAACCACAAAUACAACAAGA         57         antisense         3046   FLT1:3358L21 siRNA           3501         UUACGGAGUGUGGGAAAACACAAGA         67         sense         30778   FLT1:4753U21 siRNA           4773         UAGAGGGCCUAAGGAGAAAGGAAAGGA         67         sense         30778   FLT1:4753U21	FLT1		1	22	antisense	30340	Sense is caps FLT1:2358L21 siRNA	uuGuuGuAuruuGuGGuuGTX	423
1 2338         AACAACCACAAAUACAACAAGA         57 antisense         30342 EL71:2358L21 siRNA           2 2338         AACAACCACAAAUACAACAAGA         57 antisense         30343 FLT1:2358L21 siRNA           2 2338         AACAACCACAAAAUACAACAAGA         57 antisense         30345 FLT1:2358L21 siRNA           2 2338         AACAACCACAAAUACAACAAGA         57 antisense         30345 FLT1:2358L21 siRNA           2 2338         AACAACCACAAAUACAACAAGA         57 antisense         30346 FLT1:2358L21 siRNA           2 2338         AACAACCACAAAUACAACAAGA         57 antisense         30346 FLT1:2358L21 siRNA           2 2338         AACAACCACAAAUACAACAAGA         57 antisense         30446 FLT1:2358L21 siRNA           2 2338         AACAACCACAAAUACAACAAGA         57 antisense         30776 FLT1:2358L21 siRNA           2 2338         AACAACCACAAAUACAACAAGA         57 antisense         30777 FLT1:134U21 siRNA           2 3501         UUACGGAGUUGGACCAUCAU         60 sense         30778 FLT1:4753021 siRNA           3 4751         AGCAAAAGCAAGGAGAAAGA         62 sense         30779 FLT1:4753021 siRNA           4 4751         AGCAAAAGCAAGGAGAAAGA         63 antisense         30780 FLT1:4771021 siRNA           4 4751         UAGCAGGCCUAAGAAAGGAGAAAGA         61 antisense         30780 FLT1:477112 siRNA           4 4751	FLT1		1	57	antisense	30341	FLT1:2358L21 siRNA	uuGuuGuAuuuuGuGGuuGTX	424
2338         AACAACCACAAAAUACAACAAGA         57         antisense         30343 FLT1:2358L21 siRNA           2338         AACAACCACAAAAUACAACAAGA         57         antisense         30344 FLT1:2358L21 siRNA           2338         AACAACCACAAAAUACAACAAGA         57         antisense         30346 FLT1:2358L21 siRNA           2338         AACAACCACAAAUACAACAAGA         57         antisense         30416 FLT1:2358L21 siRNA           2338         AACAACCACAAAUACAACAAGA         57         antisense         30416 FLT1:2358L21 siRNA           2338         AACAACCACAAAUACAACAAGA         57         antisense         30717 FLT1:184U21 siRNA           2338         AACAACCACAAAUACAACAAGA         61         sense         30776 FLT1:4718U21 siRNA           3501         UUACGGAGUAUUGCUGUGGAAA         61         sense         30778 FLT1:4715U21 siRNA           4751         AGCAAAAAGGAAAGGA         63         sense         30778 FLT1:4715U21 siRNA           4751         AGCAAAAAGGAAAGGAGGAGAAAAGA         63         sense         30778 FLT1:4732U21 siRNA           4751         AGCAAAAAGGAGGAGAAAAGA         63         sense         30780 FLT1:4732U21 siRNA           4751         UAGCGGGCCUAAGACAGGAGAAAAGA         61         antisense         30781 FLT1:4771L21 siRNA	FLT1			22	antisense	30342	FLT1:2358L21 siRNA	uuGuuGuAuuuuGuGGuuGTU	425
2338         AACAACCACAAAAUACAACAAGA         57         antisense         30344 FLT1:2358L21 siRNA           2338         AACAACCACAAAAUACAACAAGA         57         antisense         30346 FLT1:2358L21 siRNA           2338         AACAACCACAAAAUACAACAAGA         57         antisense         30346 FLT1:2358L21 siRNA           2338         AACAACCACAAAAUACAACAAGA         57         antisense         30446 FLT1:2358L21 siRNA           1182         UCGUGUAAGGAGACACAAGA         57         antisense         30777 FLT1:134U21 siRNA           3501         UUACGGAGUAUUGCUGUGGGAAA         61         sense         30777 FLT1:144U21 siRNA           4713         UAGCAGGCCUAAGACAUGUGGGAAA         62         sense         30778 FLT1:4753U21 siRNA           4713         UAGCAGGCCUAAGACAUGUGGGGAAA         61         antisense         30780 FLT1:4753U21 siRNA           4713         UAGCAGGCCUAAGACAUGUGGGAAA         61         antisense         30781 FLT1:4753U21 siRNA           4771         UACGGAGUAUUGCUGGGAAAAGG         61         antisense         30781 FLT1:4753U21 siRNA           4771         UACGGAGUAUUGCUGGGAAA         61         antisense         30781 FLT1:4753U21 siRNA           4771         AGCAAAAAGCAUGUGGGAAAAGG         62         antisense         30781 FLT1:4753U21 siRNA	FLT1		AACAACCACAAAAUACA	22	antisense		FLT1:2358L21 siRNA	uuGuuGuAuuuuGuGGuuGTt	426
2338         AACAACCACAAAUACAACAAGA         57         antisense         30345 FLT1:2358L21 siRNA           2338         AACAACCACAAAUACAACAAGA         57         antisense         30346 FLT1:2358L21 siRNA           2338         AACAACCACAAAUACAACAAGA         57         antisense         30416 FLT1:2358L21 siRNA           1182         UCGUGUAAGGAGUGGACCAUCAU         60         sense         30777 FLT1:1184U21 siRNA           3501         UUACGGAGUAUUGCUGGGAAA         61         sense         30777 FLT1:1184U21 siRNA           4713         UAGCAGGCCUAAGACAUGUGGAAAGG         62         sense         30778 FLT1:4503U21 siRNA           4751         AGCAAAAGGAGGGAGAAAGG         63         sense         30778 FLT1:4753U21 siRNA           4751         AGCAAAAGGAGGGAGAAAGG         63         antisense         30780 FLT1:3521L21 siRNA           4751         AGCAAAAGGAGGAGAAAGG         63         antisense         30781 FLT1:3521L21 siRNA           4713         UAGCAGAGGAGAAAGGAAAGG         62         antisense         30781 FLT1:4733L21 siRNA           4751         AGCAAAAAGGAAAAGGAAAAGGAAAAGG         62         antisense         30781 FLT1:471L121 siRNA           4751         AGCAAAAAGGAAAAGGAAAAAGGAAAAGGAAAAGGAAAAGGAAAA	FLT1	<del>                                     </del>	L	22	antisense	30344	FLT1:2358L21	uuGuuGuAuuuuGuGGuuGTu	427
2338         AACAACCACAAAUACAACAAGA         57         antisense         30346         FLT1:2358L21 siRNA           2338         AACAACCACAAAUACAACAAGA         57         antisense         30416         FLT1:2358L21 siRNA           1182         UCGUGUAAGGAGUGGACCAUCAU         60         sense         30777         FLT1:144U21 siRNA           3501         UUACGGAGUAGGACAUGUGGGAAA         61         sense         30777         FLT1:4753U21 siRNA           4713         UAGCAGGCCUAAGACAUGUGAGG         62         sense         30778         FLT1:4753U21 siRNA           4751         AGCAAAAAGCAAGGGAAAAGA         63         sense         30780         FLT1:4753U21 siRNA           4751         AGCAAAAAGCAAGGGAGAAAAGA         63         sense         30780         FLT1:4753U21 siRNA           4751         AGCAAAAAGCAAGGAGAGAAAAGA         61         antisense         30781         FLT1:4753U21 siRNA           4713         UAGCAGGCCUAAGGAGAAAAGA         61         antisense         30782         FLT1:4733L21 siRNA           4751         AGCAAAAAGCAAGGAGAGAGAAAAGA         62         antisense         30782         FLT1:4733L21 siRNA           4751         AGCAAAAAGCAAGGAGAGAGAGAGAGAGAGAGAGAGAGA	FLT1		L	22	antisense		FLT1:2358L21 siRNA (2340C) idT	uuGuuGuAuuuuGuGGuuGTD	428
2338         AACAACCACAAAUACAACAAGA         57         antisense         30416         FL71:2358L21 siRNA           1182         UCGUGUAAGGAGUGGACCAUCAU         60         sense         30777         FL71:184U21 siRNA           3501         UUACGGAGUAUUGCUGGGAAA         61         sense         30778         FL71:3503U21 siRNA           4713         UAGCAGGCCUAAGACAUGUGGGAAAAGA         62         sense         30779         FL71:4715U21 siRNA           4751         AGCAAAAAGCAAGGGAGAAAAGA         63         sense         30779         FL71:4753U21 siRNA           4751         AGCAAAAGCAAGGGAGAAAAGA         63         sense         30780         FL71:4753U21 siRNA           4751         AGCAAAAGGAGGGAGAGAAAAGA         61         antisense         30781         FL71:4733L21 siRNA           4713         UAGCAGGCCUAAGACAUGUGGGAAA         61         antisense         30782         FL71:4733L21 siRNA           4751         AGCAAAAAGCAAGGGAGAAAAGA         62         antisense         30783         FL71:4733L21 siRNA           4751         AGCAAAAAGCAAGGAGAAAAGA         63         antisense         30784         FL71:4771L21 siRNA           4751         AGCAAAAAGCAAAAGAAAAGAAAAGA         63         antisense         30784         FL71:4771L21 siRNA	FL71			57	antisense	30346	FLT1:2358L21	uuGuuGuAuuuuGuGGuuGXT	429
1182         UCGUGUAAGGAGUGGACCAUCAU         60         sense         30777         FLT1:1184U21 siRNA           3501         UUACGGAGUAUUGCUGUGGGAAA         61         sense         30779         FLT1:1184U21 siRNA           4713         UAGCAGGCCUAAGACAUGUGGGAAAGG         62         sense         30779         FLT1:4715U21 siRNA           4751         AGCAAAAAGCAAGGGAGAAAGA         63         sense         30780         FLT1:4753U21 siRNA           1182         UCGUGUAAGGAGAGAAAGA         63         antisense         30781         FLT1:4753U21 siRNA           1182         UCGUGUAAGGAGUGGACCAUCAU         60         antisense         30781         FLT1:4753L21 siRNA           3501         UUACGGAGUAUUGCUGUGGGAAA         61         antisense         30782         FLT1:4733L21 siRNA           4713         UAGCAGACAGGGAGAAAGA         62         antisense         30783         FLT1:4733L21 siRNA           4751         AGCAAAAAGCCAAGGGAGAAAGA         63         antisense         30783         FLT1:4733L21 siRNA           4751         AGCAAAAAGCCAAGGAGAAAGA         63         antisense         30784         FLT1:4733L21 siRNA           4751         AGCAAAAAGCAAGGAAAAGA         63         antisense         30784         FLT1:471L121 siRNA	FLT1			57	antisense		FLT1:2358L21 siRNA	uuGuuGuAuuuuGuGGuuGTsT	430
3501         UUACGGAGUAUUGCUGUGGGAAA         61         sense         30778         FLT1:3503U21 siRNA           4713         UAGCAGGCCUAAGACAUGUGAGG         62         sense         30779         FLT1:4715U21 siRNA           4751         AGCAAAAAGCAAGAGAAAAGA         63         sense         30779         FLT1:4715U21 siRNA           4751         AGCAAAAAGCAAGGAGAAAAGA         63         sense         30780         FLT1:4753U21 siRNA           1182         UCGUGUAAGGACAUGUGGACAUCAU         60         antisense         30781         FLT1:3521L21 siRNA           3501         UUACGGAGUAUUGCUGUGGAAA         61         antisense         30782         FLT1:3521L21 siRNA           4713         UAGCAGGCCUAAGACAUGUGAGG         62         antisense         30782         FLT1:4733L21 siRNA           4751         AGCAAAAAGCAAGGAGAAAGA         63         antisense         30782         FLT1:4771L21 siRNA           4751         AGCAAAAAGCAAAAGGAAAAGAAAAGAAAAAGAAAAAAAA	FLTA	<del> </del>	UCGUGUAAGGAGUGGACCAUCAU	09	sense	30777	FLT1:1184U21 siRNA	B GuGuAAGGAGuGGAccAucTT B	431
4713         UAGCAGGCCUAAGACAUGUGAGG         62         sense         30779         FLT1:4715U21 siRNA           4751         AGCAAAAAGCAAGGAGAAAAGA         63         sense         30780         FLT1:4715U21 siRNA           1182         UCGUGUAAGGAGUGGACCAUCAU         60         antisense         30781         FLT1:4753U21 siRNA           3501         UUACGGAGUAUUGCUGUGGAAA         61         antisense         30782         FLT1:3521L21 siRNA           4713         UAGCAGGCCUAAGACAUGUGAGG         62         antisense         30783         FLT1:4733L21 siRNA           4751         AGCAAAAAGCCAAGGGAGAAAAGA         63         antisense         30784         FLT1:4771L21 siRNA           2338         AACAACCACAAAAUACAACAAGA         57         sense         30955         FLT1:2340U21 siRNA	FLT1	<del>                                     </del>		61	sense	30778	FLT1:3503U21 siRNA	B AcGGAGuAuuGcuGuGGGGATT B	432
4751         AGCAAAAAGCAAGGGAGAAAGA         63         sense         30780         FLT1:4753U21 siRNA           1182         UCGUGUAAGGAGUGGACCAUCAU         60         antisense         30781         FLT1:1202L21 siRNA           3501         UUACGGAGUAUUGCUGUGGAAA         61         antisense         30782         FLT1:3521L21 siRNA           4713         UAGCAGGCCUAAGACAUGUGAGG         62         antisense         30783         FLT1:4733L21 siRNA           4751         AGCAAAAAGCAAGAGAAAGGA         63         antisense         30783         FLT1:4771L21 siRNA           2338         AACAACCACAAAAUACAACAAGA         57         sense         30955         FLT1:2340U21 siRNA	FLT1			62	sense		FLT1:4715U21 siRNA	B GcAGGccuAAGAcAuGuGATT B	433
1182         UCGUGUAAGGAGUGGACCAUCAU         60         antisense         30781         FLT1:1202L21 siRNA           3501         UUACGGAGUAUUGCUGUGGGAAA         61         antisense         30782         FLT1:3521L21 siRNA           4713         UAGCAGGCCUAAGACAUGUGAGG         62         antisense         30783         FLT1:4731L21 siRNA           4751         AGCAAAAAGCAAGGGAAAAGA         63         antisense         30784         FLT1:477LL21 siRNA           2338         AACAACCACAAAAUACAACAAGA         57         sense         30955         FLT1:2340U21 siRNA	FLT1			63	seuse	30780	FLT1:4753U21 siRNA	B cAAAAAGCAAGGAAAATT B	434
3501         UUACGGAGUAUUGCUGUGGGAAA         61         antisense         30782         FLT1:3521L21 siRNA           4713         UAGCAGGCCUAAGACAUGUGAGG         62         antisense         30783         FLT1:4733L21 siRNA           4751         AGCAAAAAGCAAGGGAAAAGA         63         antisense         30784         FLT1:4771L21 siRNA           2338         AACAACCACAAAAUACAACAAGA         57         sense         30955         FLT1:2340U21 siRNA	FLT1		UCGUGUAAGGAGUGGACCAUCAU	09	antisense	30781	FLT1:1202L21 siRNA '1184C) stabn5	GAuGGuccAcuccuuAcAcTsT	435
4713         UAGCAGGCCUAAGACAUGUGAGG         62         antisense         30783         FLT1:4733L7 siRNA           4751         AGCAAAAAGGAAAAGGA         63         antisense         30784         FLT1:4771L21 siRNA           2338         AACAACCACAAAAUACAACAAGA         57         sense         30955         FLT1:2340U21 siRNA	FLT1	3501	UVACGGAGUAUUGCUGUGGGAAA	61	antisense		FLT1:3521L21 siRNA 3503C) stah05	ucccAcAGcAAuAcuccGuTsT	436
4751         AGCAAAAAGCAAGGGAGAAAAGA         63         antisense         30784         FLT1:4771L21 siRNA           2338         AACAACCACAAAAUACAACAAGA         57         sense         30955         FLT1:2340U21 siRNA	FLT1	4713	UAGCAGGCCUAAGACAUG	62	antisense	30783	FLT1:4733L21 siRNA 4715C) stab05	ucAcAuGucuuAGGccuGcTsT	437
2338 AACAACCACAAAAUACAACAAGA 57 sense 30955 FLT1:2340U21 siRNA	7	4751	AGCAAAAGCAAGGGAGA	63		30784 F	FLT1:4771L21 siRNA 4753C) stab05	uuuucucccuuGcuuuuuGTsT	438
	F	2338		22		30955 F	-LT1:2340U21 siRNA	B cAccAcAAAuAcAACAATT B	439

	440	441	442	443	444	445	446	7447	448		449	9	451	452	3	453	454	455	,	450	45/	458	459	460	461	
	uuGuuGuAuuuuGuGGuuGTsT	AACAACAUAAAACACCAACTT	GUUGGUGUUUNAUGUUGUUTT	B AACAACAUAAAACACCAACTT B	GuuGGuGuuuuAuGuuGuuTsT	B AACAACAUAAAACACCAACTT B	TsTillelillelilleli		CUGAGUUUAAAAGGCACCCTT	GCAAGGAGGGCCUCUGAGGII	CCUGGAAGAAUCAAAACCII	GGGUGCCUUUAAACUCAGII	CAUCAGAGGCCCUCCUUGCTT	GGIIIIIIGAUUCUUUCCAGGTT		B cuGAGuuuAAAAGGcAcccTT B	B GcAAGGAGGCcucuGAuGTT B	B ccuGGAAAGAAucAAAAccTT B	B	GGGuGccuuuuAAAcucAG1s1	cAucAGAGGcccuccuuGcTsT	GGuuuuGAuucuuuccAGGTsT	B cuGAGuuuAAAAGGcAccTT B	B GcAAGGAGGccucuGAuGTT B	B CONSEASON B	
stah07	antisense 30956 FLT1:2358L21 siRNA	20063 El T1-03401 121 siRNA inv	30964 FLT1:2358L21 siRNA	30965 FLT1:2340U21 siRNA	30966 FLT1:2358L21 siRNA	30967 FLT1:2340U21 siRNA	stab07 inv	30968 FLT1:2358L21 siRNA (2340C) stab08 inv	31182 FLT1:349U21 siRNA TT	31183 FLT1:2949U21 siRNA TT	31184 FLT1:3912U21 siRNA TT	31185 FLT1:367L21 siRNA	antisense 31186 FLT1:2967L21 siRNA	(2949C) 11 24403 F1 74:20201 24 SIDNA	3118/ FLI 1:3930LZ I SINNA (3912C) TT	31188 FLT1:349U21 siRNA	31189 FLT1:2949U21 siRNA	Stab04 24400 Et T4:30421124 siRNA	31190 FL11.3812021 31137   stab04	31191 FLT1:367L21 siRNA (1349C) stab05	31192 FLT1:2967L21 siRNA	antisense 31193 FLT1:3930L21 siRNA	31194 FLT1:349U21 siRNA	31195 FLT1:2949U21 siRNA	stab07	31196 FLT1:3912U21 SIKINA stab07
	antisense 30	十	antisense 3	sense 3	antisense 3	sense 3	-	antisense 3	sense 3	sense 3	1	antisense 3	antisense 3		antisense	sense	sense	-	seuse	antisense (	antisense	antisense	sense	sense		sense
-	27	12	21 2	22	22	57		22	26	29	88	56	59	1	28	56	59	5	8	26	29	28	26	55	}	28
	AACAACCACAAAAUACAACAAGA		AACAACCACAAAAUACAACAAGA	AACAACCACAAAAUACAACAAGA	AACAACCACAAAAUACAACAAGA	AACAACCACAAAAIIACAACAAGA		AACAACCACAAAAUACAACAAGA	AACLIGAGUUUAAAAGGCACCCAG		AGCCIIGGAAAGAAUCAAAACCUU	AACUGAGUUUAAAAGGCACCCAG	AAGCAAGGAGGCCUCUGAUGGU		AGCCUGGAAAGAAUCAAAACCUU	AACUGAGUUUAAAAGGCACCCAG	AAGCAAGGAGGGCCUCUGAUGGU		AGCCUGGAAAGAAUCAAAACCUU	AACUGAGUUUAAAAGGCACCCAG	AAGCAAGGAGGCCUCUGAUGGU	AGCCUGGAAAGAAUCAAAACCUU	AACUGAGUUUAAAAGGCACCCAG	- 1		AGCCUGGAAAGAAUCAAAACCUU
-	2338		2338	2338	2338	2238	2000	2338	347	$\top$	3010	347	2947		3910	347	2947		3910	347	2947	3910	347	-+	7847	3910
	FLT1		FLT1	FLT1	FLT1	14		FLT1	El T1	1	- 1-1-1	FLT1	FLT1		FLT1	FLT1	17		FLT1	FLT1	FLT1	FLT1	F	i	7	FLT1

462	463	464	465	466	467	468	469	470	471	472	473	474	475	476	477	478	479	480	481	482
GGGuGccuuuuAAAcucAGTsT	cAucAGAGGcccuccuuGcTsT	GGuuuuGAuucuuuccAGGTsT	CCCACGGAAAAUUUGAGUCTT	GUAGUCUCCGGGAGGAACGTT	CCAAAACUAAGAAAGGUCCTT	GACUCAAAUUUUCCGUGGGTT	CGUUCCUCCGGAGACUACTT	GGACCUUUCUUAGUUUUGGTT	B cccAcGGAAAAuuuGAGucTT B	B GuAGucuccGGGAGGAAcGTT B	B ccAAAAcuAAGAAAGGuccTT B	GAcucAAAuuuuccGuGGGTsT	cGuuccuccGGAGAcuAcTsT	GGAccunucuuAGuuuuGGTsT	B cccAcGGAAAAuuuGAGucTT B	B GuAGucuccGGGAGGAAcGTT B	B ccAAAAcuAAGAAAGGuccTT B	GAcucAAAuuuuccGuGGGTsT	cGuuccuccGGAGAcuAcTsT	GGAccuuucuuAGuuuuGGTsT
antisense 31197 FLT1:367L21 siRNA (349C) stab08	31198 FLT1:2967L21 siRNA (2949C) stab08	31199 FLT1:3930L21 siRNA (3912C) stab08	31200 FLT1:349U21 siRNA inv	31201 FLT1:2949U21 siRNA inv TT	31202 FLT1:3912U21 siRNA inv TT	31203 FLT1:367L21 siRNA (349C) inv TT	31204 FLT1:2967L21 siRNA (2949C) inv TT	31205 FLT1:3930L21 siRNA (3912C) inv TT	31206 FLT1:349U21 siRNA stab04 inv	31207 FLT1:2949U21 siRNA stab04 inv	31208 FLT1:3912U21 siRNA stab04 inv	31209 FLT1:367L21 siRNA (349C) stab05 inv	31210 FLT1:2967L21 siRNA (2949C) stab05 inv	31211 FLT1:3930L21 siRNA (3912C) stab05 inv	31212 FLT1:349U21 siRNA stab07 inv	31213 FLT1:2949U21 siRNA stab07 inv	31214 FLT1:3912U21 siRNA stab07 inv	31215 FLT1:367L21 siRNA (349C) stab08 inv	31216 FLT1:2967L21 siRNA (2949C) stab08 inv	
antisense 37	antisense 37	antisense 37	sense 3.	sense 3.	sense 3	antisense 37	antisense 37	antisense 31	sense 31	sense 31	sense 31	antisense 31	antisense 31	antisense 31	sense 31	sense 31	sense 31	antisense 31	antisense 31	antisense 31217
99	29	28	26	59	28	26	29	58	56	59	58	56	59	58	56	59	58	56	59	58
AACUGAGUUUAAAAGGCACCCAG	_	ſ.	AACUGAGUUUAAAAGGCACCCAG	AAGCAAGGAGGCCUCUGAUGGU	AGCCUGGAAAGAAUCAAAACCUU	AACUGAGUUUAAAAGGCACCCAG	AAGCAAGGAGGCCUCUGAUGGU	AGCCUGGAAAGAAUCAAAACCUU	AACUGAGUUUAAAAGGCACCCAG	AAGCAAGGAGGCCUCUGAUGGU	AGCCUGGAAAGAAUCAAAACCUU	AACUGAGUUUAAAAGGCACCCAG	AAGCAAGGAGGCCUCUGAUGGU	AGCCUGGAAAGAAUCAAAACCUU	AACUGAGUUUAAAAGGCACCCAG	AAGCAAGGAGGCCUCUGAUGGU	AGCCUGGAAAGAAUCAAAACCUU	AACUGAGUUUAAAAGGCACCCAG	AAGCAAGGAGGCCUCUGAUGGU	AGCCUGGAAAGAAUCAAAACCUU
347	2947	3910	347	2947	3910	347	2947	3910	347	2947	3910	347	2947	3910	347	2947	3910	347	2947	3910
FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	PLT1	FLT1	FLT1

PCT/US03/05028

WO 03/074654

483	484	485	486	9	487	488	489	490	491	492	703	Op 1	494	495	496	107	9	498	499	200	501	3	205	503	504
B CUGAGUUUAAAAGGCACCCTT B	B GCAAGGAGGCCUCUGAUGTT B	B CCUGGAAAGAAUCAAAACCTT B	ToTOVOLOVA ALII II	GGGUGCCUUUUAAACUCAGISI	CAUCAGAGGCCCUCCUUGCTsT	GGUUUUGAUUCUUUCCAGGTsT	B CCCACGGAAAAUUUGAGUCTT B	B GUAGUCUCCGGGAGGAACGTT B	B CCAAAACUAAGAAAGGUCCTT B	GACTICAAAIIJIJOCCGUGGGTST	To+C v 110 v 0 v 0 v 0 v 0 v 0 v 0 v 0 v 0 v	CGUUCCUCCGGAGACUACISI	GGACCUUUCUUAGUUUUGGTST	unGunGnAnnnuGnGGnuGXsX	XeXaBlugarianaOAAA	CAUCAGAGGCCCGCCGGCGGC	uuGuuGuAuuuuGuGGuuGXs1	cAucAGAGGcccuccuuGcXsT	B CAACCACAAAAUACAACAATT B	B AACAACAUAAAACACCAACTT B	Tatol III Golden III III III III III III III III III I	UUGUUGUANAAAAAAAAA	GUUGGUGUUUAUGUUGUUTST	B cAAcuGAGAAGccAAGAcuTT B	B cAuGGAccuAucuGGGuccTT B
31270	31271	07.040	31272	31273	31274	31275	31276	31277	31278		31279	antisense 31280 FLT1:2967L21 siRNA	31281	31424		ise 31425 FLT1:2967L21 siRNA (2949C) stab11 3'-BrdU	31442	31443	31449	stab09 31750 Ft T1-2340[121 siRNA inv	24.5	nse 31451 FLT1:2358L21 siRNA	31452	(2340C) IIIV Stab 10	
sense	seuse	-	seuse	antisense	antisense	antisense	sense	seuse	Sense		6 antisense	+	8 antisense	7 antisense		g antisense	7 antisense	59 antisense	57 sense		esuse /c	57 antisense	57 antisense	$\dashv$	64 sense 65 sense
A A CHARLEM IN TABABAGG LACTORG   56			AGCCUGGAAAGAAUCAAAACCUU 58	AACUGAGUUUAAAAGGCACCCAG 56	- 1	AGCCI IGGAAAGAAUCAA				AGCCOGGAAAGAAGCAAAGGAA	AACUGAGUUUAAAAGGCACCCAG 56	AAGCAAGGAGGCCUCUGAUGGU 59		- 1	AACAACCACAAAAACACAACA	AAGCAAGGAGGCCUCUGAUGGU 59	AACAACCACAAAAUACAACAAGA 57	AAACCAAAGGGCCUCUGAUGGU	AACAACCACAAAAIIACAACAAGA	AACHACCACCACCACCACCACCACCACCACCACCACCACC	AACAACCACAAAAUACAACAAGA	AACAACCACAAAAUACAACAAGA	AACAACCACAAAAUACAACAAGA		AGCAACUGAGAAGCCAAGACUGA AGCAACUGAACCIMICIFEGIICUII
247	3047	FLI1 294/	FLT1 3910	FLT1 347	2047				$\dashv$	FLT1 3910	FLT1 347	FI T1 2947	-		FLT1   2338	FLT1 2947	FI T1 2338	-		FL11 2338	FLT1 2338	FI T1 2338			FOS 17

•		

UAGGGAGGACCUUAUCUGUGCGU 66 sense 30771 FOS:1405U21 siRNA
67
or sense ourre
AGCAACUGAGAAGCCAAGACUGA 64 antisense 30773 FOS:37L21 siRNA (19C) stab05
IUCCUU 65 antisense 30774 FOS:1046L21 siRNA (1028C) stab05
UAGGGAGGACCUUAUCUGUGCGU 66 antisense 30775 FOS:1423L21 siRNA (1405C) stab05
AAGCAUCCAUGUGUGGACUCAAG 67 antisense 30776 FOS:1480L21 siRNA (1462C) stab05
AGCAACUGAGAAGCCAAGACUGA 64 sense 31049 FOS:19U21 siRNA
esues gense
1403 UAGGGAGGACCUUAUCUGUGCGU 66 sense 31051 FOS:1405U21 siRNA
67 sense
AGCAACUGAGAAGCCAAGACUGA 64 antisense 31125 FOS:37L21 siRNA (19C)
GACAUGGACCUÁUCUGGGUCCUU 65 antisense 31126 FOS:1046L21 siRNA
UAGGGAGGACCUUAUCUGUGCGU 66 antisense 31127 FOS:1423L21 siRNA (1405C)
AAGCAUCCAUGUGUGGACUCAAG 67 antisense 31128 FOS:1480L21 siRNA (1462C)
UGAAGAGGGAAAGCUGACAUCUG 68 sense 31541 GAB2:2681U21 siRNA
GAGGAAGAAGGAAGGAGGCUU 69 sense 31542 GAB2:4316U21 siRNA
GAGAGGACUGAGCCUACGGAAAG 70 sense 31543 GAB2:5006U21 siRNA
UUUGCUGUGGUGACACAUGGUAC 71 sense 31544 GAB2:5958U21 siRNA
UGAAGAGGGAAAGCUGACAUCUG 68 antisense 31545 GAB2:2699L21 siRNA (2681C)
GAGGAAGAAGGAAGGCUU 69 antisense 31546 GAB2:4334L21 siRNA (4316C)
GAAAG 70 antisense 31547 GAB2:5024L21 siRNA
GGUAC 71 antisense 31548 GAB2:5976L21 siRNA
72 antisense 25245 RPI 17763 Her2Neu AS as siRNA Str 2 (antisense)
73 sense 25246 RPI 17763 Her2Neu AS as siRNA Str 1 (sense)
72 sense 25247 RPI 17763 Her2Neu AS

	530	531	532	533	534	535	536	537	538	539	540	541	542	543	544	545	546	547	548	549
	B UCGGCGUCACUCGUGGUACCU B	uccaugeugeucacugegeeuuu	AGCCGCAGUGAGCACCAUGGAUU	B UCCAUGGUGCUCACUGCGGCUUU B	B AGCCGCAGUGAGCACCAUGGAUU B	UGGGGUCGUCAAAGACGUUTT	AACGUCUUUGACGACCCCATT	UUGCAGAAACUGCUGGGGUTT	ACCCCAGCAGUUUCUGCAATT	GGUGCUUGGAUCUGGCGCUTT	AGCGCCAGAUCCAAGCACCTT	UCGCGGUCUAGGUUCGUGGTT	CCACGAACCUAGACCGCGATT	GAUCUUUGGGAGCCUGGCATT	UGCCAGGCUCCCAAAGAUCTT	ACGGUCCGAGGGUUUCUAGTT	CUAGAAACCCUCGGACCGUTT	GsGsusGscuuGGAucuGGcGscsusTsT	AsGsCsGsCsCAGAUCCAAGCACCTsT	GSGSUSGSCSUUGGAUCUGGCGCUTST
as siRNA Str 1 (sense) Inverted control	25248 RPI 17763 Her2Neu AS as siRNA Str 1 (sense) Inverted control compliment	RPI 17763 Her2Neu AS as siRNA Str 2 (antisense)+2U overhand	25823 RPI 17763 Her2Neu AS as siRNA Str 1 (sense)+2U overhang	antisense 25842 RPI 17763 Her2Neu AS as siRNA Str 2 (antisense)+2U overhang	25843 RPI 17763 Her2Neu AS as siRNA Str 1 (sense)+2U overhang	28262 Her2.1.sense Str1	28263 Her2.1.antisense Str2	28264 Her2.1.sense Str1 inverted	28265 Her2.1.antisense Str2 inverted	28266 Her2.2.sense Str1	28267 Her2.2.antisense Str2	28268 Her2.2.sense Str1 inverted	28269 Her2.2,antisense Str2 inverted	28270 Her2.3.sense Str1	28271 Her2.3.antisense Str2	28272 Her2.3.sense Str1 inverted	28273 Her2.3.antisense Str2 inverted	29989 Her2.2.sense Str1 (site 2344)	29990 Her2.2.antisense Str2	29991 Her2.2.sense Str1 (site 2344)
	25248		25823	25842	25843	28262	28263	28264	28265	28266				28270	-	28272	28273	29989		
	sense	antisense 25822	sense	antisense	sense	sense	antisense	seuse	antisense	seuse	antisense	sense	antisense	seuse	antisense	seuse	antisense	sense	antisense	seuse
	74	72	73	72	73	75	75	75	75	92	92	92	92	77	77	2.2	77	76	9/	76
	CAUGGUGCUCACUGCGGCU	CCGCAGUGAGCACCAUGGA	AGCCGCAGUGAGCACCAUG	CCGCAGUGAGCACCAUGGA	AGCCGCAGUGAGCACCAUG	UGGGGUCGUCAAAGACGUU	UGGGGUCGUCAAAGACGUU	UGGGGUCGUCAAAGACGUU	UGGGGUCGUCAAAGACGUU	GGUGCUMGGAUCUGGCGCU	GGUIGGAUCUGGCGCU	GGUGCUUGGAUCUGGCGCU	GEUGCUUGGAUCUGGCGCU	GAUCUUUGGGAGCCUGGCA	GAUCUUUGGGAGCCUGGCA	GAUCUUUGGGAGCCUGGCA	GAUCUUUGGGAGCCUGGCA	GGUGCUUGGAUCUGGCGCU	GGUGGAUCUGGCGCU	GEUGCUUGGAUCUGGCGCU
							3706		3706		2344		2344					2342	2344	2342
	Her2	Her2	Her2	Her2	Her2	Her2	Her?	Her2	Her2	Hor	Fers	Her2	Her2	Hero	Hero	Her2	Her2	Her2	Hor	Her2

o	2
	٦
_	

571	572	573	574	575	576	577	565	578	579	580	566	581	582	583	584	700	- oc	562	282	3 5	28/	000	SOC -
B GGuGcuuGGAucuGGcGcuTT B	BucGcGGucuAGGuucGuGGTT B	B GGuGcuuGGAucuGGcGcuTT B	B AAcGucuuuGAcGAcccATT B	AGcGccAGAuccAAGcAccTsT	uGGGGucGucAAAGAcGuuTsT	B GAAuGGcucAGuGAccuGuTT B	B GGuGcuuGGAucuGGcGcuTT B	B AAcGucuuuGAcGAcccATT B	B cAccuucAAAGGGAcAccuTT B	AcAGGucAcuGAGccAuucTsT	AGcGccAGAuccAAGcAccTsT	uGGGGucGucAAAGAcGuuTsT	AGGuGuccunuGAAGGuGTsT	B uGGGGucGucAAAGAcGuuTT B	TaTA222AG2AG2	AACGUCUUUGAKGAACGAAGA	B uGGGGucGucAAAGAcGuuli B	AAcGucuuuGAcGAcccATsT	ACCAUUUUGUGGACGAAUATT	CUGUUGGACAUCCUGGAUALI	GGAUGCCUUCUACACGUUGTT	GAACCCUCCUGAUGAGAGUII	UAUUCGUCCACAAAAUGGUII
30448 Her2 sense (site 2344)	30449 Her2 sense inverted (site	30645 HER2:2346U21 siRNA	30646 HER2:3726L21 siRNA		30648 HER2:3708U21 siRNA	37 HER2:1884U21 siRNA stab04	98 HER2:2346U21 siRNA stah04	30699 HER2:3726L21 siRNA	30700 HER2:3879U21 siRNA		30702 HER2:2364L21 siRNA	03 HER2:3708U21 siRNA		(3879C) stab05 54 HFR2:37081121 siRNA	stab07	52   HER2:3726L21 siRNA  (3708C) stab08		30954 HER2:3726L21 siRNA (3708C) stab05	31525 HRAS:77U21 siRNA	31526 HRAS:154U21 siRNA	31527 HRAS:459U21 siRNA	31528 HRAS:513U21 siRNA	31529 HRAS:95L21 siRNA (77C)
30448	3044	3064		30647	3064	30697	30698			e 30701		30703	e 30704	20054		se 30952	30953		$\top$			$\top$	315
seuse	sense	sense	antisense	antisense	sense	sense	seuse	antisense	sense	antisense	antisense	sense	antisense		seuse	antisense	seuse	antisense	sense	Sense	sense	seuse	antisense
92	92	76	75	9/	75	78	9/	75	79	78	9/	75	79	75	(2)	75	75	75	8	2,	2 8	83	8
GGUGCUUGGAUCUGGCGCU	GGUGCUUGGAUCUGGCGCU	GEUGCUUGGAUCUGGCGCU	UGGGGUCGUCAAAGACGUU	GGUGCUUGGAUCUGGCGCU	UGGGGUCGUCAAAGACGUU	GAAUGGCUCAGUGACCUGU	GGUGCUUGGAUCUGGCGCU	UGGGGUCGUCAAAGACGUU	CACCUUCAAAGGGACACCU	GAAUGGCUCAGUGACCUGU	GGUGCUUGGAUCUGGCGCU	UGGGGUCGUCAAAGACGUU	CACCUUCAAAGGGACACCU		UGGGGUCGUCAAAGACGUU	UGGGGUCGUCAAAGACGUU	UGGGGUCGUCAAAGACGUU	UGGGGUCGUCAAAGACGUU	GAACCALII II III III II II II II II II II II I	OPPLIANCE CONTROLLING TO THE PROPERTY OF THE P	GCCUGUUGGACAUCCUGGADACC	GAGGAUGCCOOCOACACCOCCC	GAACCAUUUGUGGACGAAUACG
2342	2342	2344	3706	2344	3706	1882	2344	3706	3877	1882	2344	3706	3877	3	3706	3706	3706	3706	1	4		-	95
Her2	Her2	Her2	Her2	Her2	Her2	Her2	Her2	Her2	Her2	Her2	Her2	Har	T G	2	Her2	Her2	Her2	Her2	2	OF SE	HRAS	HKAS	HRAS

290	591	592	593	594	595	596	597	598	299	009	601	602	603	604	605	909	209	809	609	610	611	612	613
UAUCCAGGAUGUCCAACAGTT	CAACGUGUAGAAGGCAUCCTT	ACUCUCAUCAGGAGGGUUCTT	AGCUUGGCCAAUCCGUGCGGU	UUGCGGAGGGUGGGCCUGGGA	CUGCCGCCUUCCACCGUUCAU	ACCCACUGCCACCGCGAAGAG	GCGCGCGAUUCCCUGAGCUG	CGCACGGAUUGGCCAAGCUGA	CCAGGCCCACCCUCCGCAACC	GAACGGUGGAAGGCGGCAGGC	cunceceeneecaeneeenec	GCUCAGGGAAUCGCGCCGCGC	B ucccuunAuAAGccGAcucTT B	B uuccAccGuucAuucuAGATT B	B ccAccGuucAuucuAGAGcTT B	B GAAGAGuuGGGcucuGucATT B	GAGucGGcuuAuAAAGGGATsT	ucuAGAAuGAAcGGuGGAATsT	GcucuAGAAuGAAcGGuGGTsT	uGAcAGAGcccAAcucuucTsT	B GAAGAGccAAcuGuGuGAGTT B	B AGGGAGGAGAGGAGUUCCTT B	B GGAGUACAGCAAACUGAAGTT B
antisense   31530   HRAS:172L21 siRNA   (154C)	31531 HRAS:477L21 siRNA (459C)	31532 HRAS:531L21 siRNA (513C)	29950 hTR:33U21 siRNA	29951 hTR:101U21 siRNA	29952 hTR:235U21 siRNA	29953 hTR:382U21 siRNA	29954 hTR:494U21 siRNA	29955 hTR:53L21 siRNA (33C)	29956 hTR:121L21 siRNA (101C)	29957 hTR:255L21 siRNA (235C)	29958 hTR:402L21 siRNA (382C)	29959 hTR:514L21 siRNA (494C)	30913 hTR:64U21 siRNA stab04	30914 hTR:243U21 siRNA stab04	30915 hTR:245U21 siRNA stab04	30916 hTR:397U21 siRNA stab04	30917 hTR:82L21 siRNA (64C) stab05	30918 hTR:261L21 siRNA (243C) stab05	30919 hTR:263L21 siRNA (245C) stab05	30920 hTR:415L21 siRNA (397C) stab05	30801 IKKg:166U21 siRNA stab04	30802 IKKg:407U21 siRNA stab04	30803 IKKg:1162U21 siRNA
ntisense 31	antisense 318	antisense 318	sense 296	sense 290	sense 296	sense 296	sense 299	antisense 299	antisense 299	antisense 296	antisense 299	antisense 299	sense 306	sense 306	sense 306	sense 308	antisense 309	antisense 309	antisense 309	antisense 309	sense 308	sense 308	sense 308
81 a	82 a	83 a	84	82	98	87	88	84 a	85 a	86 'a	87 a	88 a	89	06	91	92	89 aı	90 a	91 aı	92 aı	93	94	95
GCCUGUUGGACAUCCUGGAUACC	GAGGAUGCCUUCUACACGUUGGU	cugaacccuccugaugagagugg	UCAGCUUGGCCAAUCCGUGCGGU	GGUUGCGGAGGGUGGGCCUGGGA	GCCUGCCGCCUCCACCGUUCAU	GCACCCACUGCCACCGCGAAGAG	GCGCGCGCGAUUCCCUGAGCUG	UCAGCUUGGCCAAUCCGUGCGGU	GGUUGCGGAGGGUGGGCCUGGGA	GCCUGCCGCCUCCACCGUUCAU	GCACCCACUGCCACCGCGAAGAG	GCGCGCGCGAUUCCCUGAGCUG	GCUCCCUUNANAAGCCGACUCGC	ccuuccacceuucauucuagagc	UUCCACCGUUCAUUCUAGAGCAA	GCGAAGAGUUGGGCUCUGUCAGC	GCUCCCUUNANAAGCCGACUCGC	ccuuccacceuucauucuaeaec	UUCCACCGUUCAUUCUAGAGCAA	GCGAAGAGUUGGGCUCUGUCAGC	UGGAAGAGCCAACUGUGUGAGAU	AGAGGGAGGAGAGGAGUUCCUC	AGGGAGUACAGCAAACUGAAGGC
172	477	531	31	66	233	380	492	31	66	233	380	492	62	241	243	395	62	241	243	395	166	407	1162
HRAS	HRAS	HRAS	hTR	hTR	hTR	hTR	hTR	hTR	hTR	hTR	hTR	hTR	hTR	hTR	hTR	hTR	hTR	hTR	hTR	hTR	IKKg	IKKg	IKKg

^	_	5	
ī	7	ζ	
`	٠.	•	

614	615	616	617	618	619	620	621	622	020	624	625	808	070	627	628	629	630	631	632	633	634	635	636	637	3	638	639	640	
B cAuGGAGuGcAuuGAGuAGTT B	cucAcAcAGuuGGcucuucTsT	GGAAcuccuucuccucccuTsT	cuucAGuuuGcuGuAcuccTsT	cuAcucAAuGcAcuccAuGTsT	B AccucAcuccuGccAcAATT B	B cuccuGucuuGcAuuGcAcTT B		B cAcAGcuAcAAcuGGAGCATT B	uuGuGGcAGGAGuuGAGGu181	GuGcAAuGcAAGAcAGGAGTsT	G. G	+ He Connection	uGcuccAGuuGuAGcuGuG181	ACCIICAACUCCUGCCACAATT	CITCHIGHICHUGCAUUGCACTT	HIGGACUUGUCACAACAGTT	CACAGCIJACAACUGGAGCATT	UJGUGGCAGGAGUUGAGGUTT	GLIGCAAUGCAAGACAGGAGTT	CHGINIGUGACAAGUGCAATT	HECHICCAGUUGUAGCUGUGTT	B uccAcuuAccuGAGGAGCATT B	B TTOWNEGAAGAGAUUCTT B	B UGAGGAUGGAAGA	B GGuucuuGccucAGAAGAGII B	B uGAAGGcucAAAccAGAcATT B	TSTAGGIBAGGIGGATST	TaTAnia Cirkania	GAAuccucuncchuccac
30804 IKKg:1390U21 siRNA	stab04 30805   KKg:184L21 siRNA	(166C) stab05 IKKg:425L21 siRNA	(407C) stab05 (407C) stab05 30807 IKKq:1180L21 siRNA	(1162C) stab05	(1390C) stab05	30809 IL2:30UZ1 SIRINA SIADO4	30810 ILZ:63UZ1 SIRINA SIABOT	30811 ILZ:88UZ1 SIRINA Stab04	30813 IL2:48L21 siRNA (30C)	stab05	stab05	30815 IL2:106L21 siRNA (88C)	30816 IL2:163L21 siRNA (145C)	stab05	31400 IL2:30U21 siRNA		02 IL 2:88UZ1 SIRINA	31403 IL2:145U21 SIRNA	31404 IL2:48L21 SIRINA (30C)	31405 IL2:81L21 SIRNA (63C)	antisense 31406 IL2:106L21 sIRNA (00C)	antisense   31407   L2:163L21 siRNA (1430)	stab04	30786 KDR:3854U21 siRNA	30787 KDR:4089U21 siRNA	stab04	30788 KUK:4191041 Sirvin	30789 KDR:3094L21 siRNA (3076C) stab05	antisense 30790 KDR:3872L21 sIRNA
30804	30805	30806	30807	20808	2000	30806	384	3081			3000	3081			_	3140				e 314(	314(	314(		1	$\dashv$			se 307	se 30
esues	anticense	o de	diffuscilso	arinsansa	antisetise	sense	sense	seuse	sense		antisense	antisense	antisense		seuse	sense	sense	sense	antisense	antisense	antisens	antisens	sense	seuse	sense		sense	antisense	
9					96 6	97	86	66	3 2	6	86	66	100	3	97	86	66,	100	97	86	ගි	90	101	102	103	3	104	101	102
000	00000			AGGGAGUACAGCAAACUGAAGGC	GUCAUGGAGUGCAUUGAGUAGGG	HAACCHCAACHCCUGCCACAAUG	AACHCCHGICUUGCAUUGCACUA		1		AACUCCUGUCUUGCAUUGCACUA	TICHHECACHUGUCACAAACAGUG		AACACAGCUACAACUGGA	STATE OF THE PROPERTY OF THE P		+	-+-	000100100001	十	+		_	2852 IIIIII GAGCALIGGAAGAGAGAUUCUG		4087 AUGGUUCUUGCCUCAGAAGAGG	4189 UCUGAAGGCUCAAACCAGACAAG	3074 UGUCCACUUACCUGAGGAGCAAG	
	1390	184	425	1180	1408	ç	3 2	5 8	143	28	61	90	8	143	_	8 8	اة ا	8	143	87	<u>ا</u> و	8 E	100				+	-	
	IKKg	IKKg	IKKg	IKKg	IKKg	=	<u>י</u>	2 2	2	112	12	=	7	12		2	2	2	2	2	2	2 2	看		 5 7	X R	KDR	KDR	<u> </u>

661

657 658 658 659 659 660 660

641

642

643 644 645 646 646

648

649

650

651 652 653 654 654

648

655

929

IIRNA							(3854C) stab05	
4189         UCUGAAGECUCAAACCAGACAAG         104         antisense         30782 (ADR-4099L21 siRNA           3872         UCUGAAGECULACCUGAGEGACAG         101         sense         31426 KOR:3076U21 siRNA           4882         UUGAGCAUGCACUGAGGACAGG         102         sense         31426 KOR:3094U21 siRNA           4882         UUUGAGCAUGCACAGCAGCAG         103         sense         31427 KOR:3094U21 siRNA           4887         AUGGUUCUUGCCUCAGACGACAGG         104         sense         31428 KOR:4191U21 siRNA           3852         UUUGAGCAUGCAUGCAGCAGAGGGUUCUG         102         antisense         31437 KOR:394U21 siRNA           3852         UUUCAGCAUGCACAGACAGAGGCU         103         antisense         31437 KOR:394U21 siRNA           3852         UUUCAGCCUCAACCAGACAGG         104         antisense         31437 KOR:394U21 siRNA           3852         UUUGAGCUCACAGCACAGACAGGAUCUG         105         sense         31437 KOR:394U21 siRNA           3854         GACACACAGGACAGGAUCUCUG         105         sense         31437 KOR:394U21 siRNA           3852         UUUGAGCAUGGACCAUCUCAUCUCU         107         sense         31437 KOR:394U21 siRNA           3854         GACACACAGGGAUUUCCUUUCCUUUCCUUUCGUUUCGAGAUCUCUCU         105         sense         31437 KOR:394U21 siRNA </td <td>KDR</td> <td>4087</td> <td>AUGGUUCUUGCCUCAGAAGAGCU</td> <td>103</td> <td></td> <td></td> <td>KDR:4107L21 siRNA (4089C) stab05</td> <td>cucuucuGAGGcAAGAAccTsT</td>	KDR	4087	AUGGUUCUUGCCUCAGAAGAGCU	103			KDR:4107L21 siRNA (4089C) stab05	cucuucuGAGGcAAGAAccTsT
3074         UGUCCACUUACCUGAGGAGCAAG         101         sense         31426         KDR:3376U21 siRNA           3822         UUUGAGCAUGGAAGAGAGAUCUG         102         sense         31427         KDR:3854U21 siRNA           4087         AUGEUUCUGAGAGAGAGACA         103         sense         31428         KDR:3854U21 siRNA           3074         UGUCAAGCCUCAACCACACAC         101         sense         31429         KDR:401012 siRNA           3074         UGUCAAGCCUCAACCAGACACAC         101         antisense         31430         KDR:407L21 siRNA           3852         UUUCAAGCCUCAACCAGACACAC         102         antisense         31431         KDR:381L21 siRNA           4087         AUGCCAUUGGAGAGACACACAGACACACACACACACACAC	KDR	4189	UCUGAAGGCUCAAACCAGACAAG	104	antisense		KDR:4209L21 siRNA (4191C) stab05	uGucuGGuuuGAGccuucATsT
3852         UUUGAGCAUGGAAGAGAGAUUCUG         102         sense         31427         KDR:3854U21 siRNA           4087         AUGGUUGUGGCUCAGAGAGAGGU         103         sense         31428         KDR:4088U21 siRNA           4189         UCUGAAGGCUCAGACGAGAGGU         104         antisense         31428         KDR:30421 siRNA           3074         UCUGAAGGCUCAAGCAGAGGU         102         antisense         31430         KDR:30721 siRNA           4087         AUGCUCCACUUGCAGAGAGAGACA         103         antisense         31431         KDR:30721 siRNA           4087         AUGCCACUUGCAGACAGACAGACACACAGACAGACAGACA	KDR	3074		101	seuse	31426	KDR:3076U21 siRNA	UCCACUUACCUGAGGAGCATT
4087         AUGGUUCUUGCCUCAGAAGAGCOU         103         sense         31428         KDR-4089U21 siRNA           4189         UCUGAAGGCUCAAACCAGACAG         104         sense         31429         KDR-4191U21 siRNA           3074         UGUCGACUUACCUGAGGACAGG         101         antisense         31431         KDR-3942L21 siRNA           3852         UUUGAGCAUGGAAGGACUCUG         102         antisense         31431         KDR-309L21 siRNA           4087         AUGGUUCUUGCCUCAGAAGAGCU         103         antisense         31432         KDR-4107L21 siRNA           4087         AUGGUUCUUGCCUCAGACGACACAG         104         antisense         31433         KDR-4209L21 siRNA           3302         UGACCUUGGAGCAUCCAGACACACACACACACACACACAC	KOR	3852	UUUGAGCAUGGAAGAGGAUUCUG	102	seuse	31427	KDR:3854U21 siRNA	UGAGCAUGGAAGAGGAUUCTT
4189         UCUGAAGGCUCAAACCAGACAAG         104         sense         31429         KDR:4191021 siRNA           3074         UGUCCACUUACCUGAGGAGCAAG         101         antisense         31430         KDR:3094L21 siRNA           3852         UUUGAGCAUGGAAGGAGUCUCAGAGGAGU         102         antisense         31431         KDR:4107L21 siRNA           4189         UCUGAAGGCUCAAAGGACAG         103         antisense         31432         KDR:4107L21 siRNA           3302         UCACCUUGAGCAUCUCAUCUGU         105         sense         31432         KDR:4209L21 siRNA           3302         UCACCUUGAGGACAUCUCAUCUGU         105         sense         31434         KDR:3304L21 siRNA           3302         UCACCUUGAGGCAUCUCAUCUGU         105         sense         31436         KDR:3304L21 siRNA           3302         UCACCUUGUUCAGGAGAGGAUUCUG         107         sense         31436         KDR:3304L21 siRNA           3302         UCACCUGUUUCAGAGGAUUCUGU         107         sense         31436         KDR:3304L21 siRNA           3302         UCACCUGUUUCCGAAGAGGAUUCUG         107         sense         31436         KDR:331L21 siRNA           3302         UUCACCUGUUUCCGAAGAGGAUUCUG         105         antisense         31436         KDR:331L21 siRNA </td <td><b>KOR</b></td> <td>4087</td> <td></td> <td>103</td> <td>seuse</td> <td>31428</td> <td>KDR:4089U21 siRNA</td> <td>GGUUCUUGCCUCAGAAGAGTT</td>	<b>KOR</b>	4087		103	seuse	31428	KDR:4089U21 siRNA	GGUUCUUGCCUCAGAAGAGTT
3074         UGUCCACUUACCUGAGGAGCAAG         101         antisense         31430         KDR:3094.21 siRNA           3852         UUUGAGCAUGGAAGAGGUUCUG         102         antisense         31431         KDR:3372L21 siRNA           4087         AUGGUUCUUGCCUCAAGAGAGCU         103         antisense         31432         KDR:3372L21 siRNA           4189         UCUGAGGCUCAAGCAGAGAG         104         antisense         31432         KDR:4209L21 siRNA           3302         UCUGAGCCUCAACCAGACAGGAUUCUG         105         sense         31436         KDR:3304U21 siRNA           3862         UUUGAGCAUGGAGGAUUCUG         105         sense         31436         KDR:3394U21 siRNA           3862         UUUGAGCAUGGAGGAUUCUG         105         sense         31436         KDR:3394U21 siRNA           3862         UUUGAGCAUGGAGGAUUCUG         107         sense         31436         KDR:332L21 siRNA           3862         UUUGAGCAUGGAGGAUUCUGG         105         antisense         31436         KDR:332L21 siRNA           3862         UUUGAGCAUGGAGGAUUCUGG         106         antisense         31431         KDR:332L21 siRNA           3862         UUUGAGCAUGGAGGAUUCUGG         107         antisense         31440         KDR:3362L21 siRNA	KDR	4189	UCUGAAGGCUCAAACCAGACAAG	104	sense	31429	KDR:4191U21 siRNA	UGAAGGCUCAAACCAGACATT
3852         UUUGAGCAUGGAAGAGGAUUCUG         102         antisense         31431         KDR:3872L21 siRNA           4087         AUGGUUCUUGCCUCAGAGAGCU         103         antisense         31432         KDR:4107L21 siRNA           4189         UCUGAAGGCUCAAACCAGACAG         104         antisense         31433         KDR:4209L21 siRNA           3802         UUCGACCUUGGAGGAUCUCUGUU         105         sense         31435         KDR:3854U21 siRNA           3802         UUCACCUUUCCUCAUGUCA         105         sense         31436         KDR:3854U21 siRNA           3802         UUCACCUUUCCUCUUUCCUCUUUCCUCUUUCUCUUUCUCUUUCCUCU	KDR	3074		101	antisense	31430	KDR:3094L21 siRNA (3076C)	UGCUCCUCAGGUAAGUGGATT
4087         AUGGULCUUGCCUCAGAAGACACOU         103         antisense         31432         KDR:4107L21 siRNA           4189         UCUGAAGGCUCAAACCAGACAGG         104         antisense         31433         KDR:3204U21 siRNA           3302         UGACCUUGGAGCAUCUCUGU         105         sense         31434         KDR:3304U21 siRNA           3852         UUUGAGCAUGGAGGAUCUCUG         105         sense         31436         KDR:3304U21 siRNA           3892         UCACCUGGUUCCGAGGAGGAUCCUGU         105         antisense         31436         KDR:3394U21 siRNA           3302         UGACCUGGAGCAUCCGAGGAGUCCUGU         107         antisense         31436         KDR:3394U21 siRNA           3302         UGACCUGGAGCAUCUCAUCUGU         105         antisense         31436         KDR:3394U21 siRNA           3302         UGACCUGGAGCAUCUCAUCUGU         105         antisense         31436         KDR:332L21 siRNA           3852         UUUCACCCGGAUCUCGAGGAUCUCGUC         102         antisense         31441         KDR:382L21 siRNA           3892         UCACCUGUUUCCUGUAUCAGUCA         107         antisense         31533         KRAS2:6262U21 siRNA           625         ACAAGACAGGGGAUUCUGCAGUUGAUCCCAGUUGAUCCCAGUUGAUCCCAGUUGAUCCCAGUUGAUCCCAGUUGAUCCCAGUUGAUCACUCGAGUUGAUCACUCGAGUUGAUCACUCGAGUUGA	KOR	3852		102		31431	KDR:3872L21 siRNA (3854C)	GAAUCCUCUUCCAUGCUCATT
4189         UCUGAAGGCUCAAACCAGACAAG         104         antisense         31433         KDR.4209L21 siRNA           3302         UGACCUUGGAGCAUCCAUCUGU         105         sense         31434         KDR:33304L21 siRNA           3852         UUUCAGCAUGGAGAGAGGAUUCUG         102         sense         31436         KDR:3854U21 siRNA           3892         UCACCUGGAGCAUCUCAUCUGU         105         sense         31436         KDR:332L21 siRNA           3892         UCACCUGGAGCAUCUCAUCUGU         105         antisense         31437         KDR:3392L21 siRNA           3852         UUUCAGCAGCAGCAGCAUCUCAUCUGU         105         antisense         31439         KDR:332L21 siRNA           3852         UUUCAGCAUGGAAGGGAUUCUGG         102         antisense         31440         KDR:3312L1 siRNA           3892         UCACCUGUUUCAUGAGGGA         106         antisense         31441         KDR:33612L1 siRNA           892         UUUCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGC	KDR	4087	AUGGUUCUUGCCUCAGAAGAGCU	103	antisense	31432	KDR:4107L21 siRNA (4089C)	CUCUUCUGAGGCAAGAACCTT
3302         UGACCUUGGAGCAUCUCAUCUGU         105         sense         31434         KDR:3364U21 siRNA           3852         UUUGAGCAUGGAAGAGAUUCUG         102         sense         31435         KDR:33854U21 siRNA           3892         UCACCUGUUUCCUGUAUGGAGGA         106         sense         31436         KDR:3384U21 siRNA           3946         GACAACACACAGCAGCAGCAUCUCUGU         107         sense         31437         KDR:3394BU21 siRNA           3920         UGACCUUGGAGCAUCUCAUCUGU         105         antisense         31437         KDR:3394BU21 siRNA           3892         UUUGAGCAUGGAGGAUCUCAUCUGU         105         antisense         31430         KDR:3312L21 siRNA           3892         UUUGAGCAUGGAGGAUUCUGG         107         antisense         31440         KDR:3912L21 siRNA           625         ACAAGACACAGCAGGAUUCAGUCA         107         antisense         31431         KDR:3912L21 siRNA           625         ACAAGACACAGCAGGAUUCAUGAUGC         108         sense         31533         KRAS2:625U21 siRNA           920         UUUCCUCGAAGUGCCAGUAUUCC         109         sense         31534         KRAS2:999U21 siRNA           920         UUUCCUCGAAGUGCCAGUAUUCGU         110         sense         31535         KRAS2:999U21 siRNA	<b>CDR</b>	4189	UCUGAAGGCUCAAACCAGACAAG	104		31433	KDR:4209L21 siRNA (4191C)	UGUCUGGUUUGAGCCUUCATI
3852         UUUGAGCAUGGAAGAGAGUUCUG         102         sense         31436         KDR:3854021 siRNA           3892         UCACCUGUUUCCUGUAUGGAGGA         106         sense         31436         KDR:3894021 siRNA           3946         GACAACACAGCAGCAGCAGCAGCAGUCUGU         107         sense         31437         KDR:3848021 siRNA           3302         UGACCUUGGAGCAUCUCAUCUGU         105         antisense         31437         KDR:3948021 siRNA           3852         UUUGAGCAUGGAGGAUCUCAUCUGU         105         antisense         31439         KDR:332121 siRNA           3892         UUUGAGCAUGGAAGAGGAUUCUG         102         antisense         31440         KDR:3912121 siRNA           3892         UCACCUGUUUCCUGUAUGAGGA         107         antisense         31441         KDR:3912121 siRNA           625         ACAAGACACAGGGUGUUGAUGAUGC         108         sense         31533         KRAS2:625U21 siRNA           920         UUUCCUCGAAGUGCCAGUAUUCC         109         sense         31534         KRAS2:920U21 siRNA           920         UUUCCUCGAAGUGCCAGUAUUCGU         110         sense         31535         KRAS2:999U21 siRNA           939         AUUUCUGGCGUUUUGGGGUUUUGGU         111         sense         31536         KRAS2:0999U21 siRNA	CDR	3302		105	seuse	31434	KDR:3304U21 siRNA	ACCUUGGAGCAUCUCAUCUTT
3892         UCACCUGUUUCCUGUAUGGAGGA         106         sense         31436         KDR:3894021 siRNA           3946         GACAACACAGGAAUCAGUCA         107         sense         31437         KDR:3948U21 siRNA           3302         UGACCUUGGAGCAUCUCAUCUGU         105         antisense         31438         KDR:3322L21 siRNA           3852         UUUGAGCAUGGAGGAUUCUCG         102         antisense         31440         KDR:3872L21 siRNA           3892         UCACCUGUUUUCCUGUAUGGAGGA         106         antisense         31440         KDR:3872L21 siRNA           3946         GACAACACAGGGAAUCAGUCA         107         antisense         31441         KDR:3872L21 siRNA           625         ACAAGACAGGGUGUUGAUGAUGA         107         antisense         31533         KRAS2:625U21 siRNA           920         UUUCCUCGAAGUCGUUGAUGC         108         sense         31533         KRAS2:920U21 siRNA           920         UUUCCUCGAAGUCCCAGUAUUCC         109         sense         31534         KRAS2:999U21 siRNA           920         UUUCCUCGAAGUCCCAGUUUUGGU         10         sense         31535         KRAS2:999U21 siRNA           920         UUUCCUCGAAGUCGAUGAUUGAU         11         sense         31536         KRAS2:099U21 siRNA	SOR.	3852	UUUGAGCAUGGAAGAGGAUUCUG	102	sense	31435	KDR:3854U21 siRNA	UGAGCAUGGAAGAGGAUUCTT
3946         GACAACAGAGAAUCAGUCA         107         sense         31437         KDR:3948U21 siRNA           3302         UGACCUUGGAGCAUCUCAUCUGU         105         antisense         31438         KDR:3322L21 siRNA           3852         UUUGAGCAUGGAGGAUUCUG         102         antisense         31439         KDR:3872L21 siRNA           3892         UCACCUGUUUCCUGUAUGGAGGA         106         antisense         31440         KDR:3912L21 siRNA           3946         GACAACACAGCAGGAAUCAGUCA         107         antisense         31441         KDR:3912L21 siRNA           625         ACAAGACACAGCAGGAAUCAGUCA         108         sense         31533         KRAS2:625U21 siRNA           920         UUUCCUCGAAGUGCCAGUAUUCC         108         sense         31533         KRAS2:625U21 siRNA           920         UUUCCUCGAAGUGCCAGUAUUCC         109         sense         31534         KRAS2:920U21 siRNA           920         UUUCCUCGAAGUGCCAGUAUUCC         109         sense         31534         KRAS2:920U21 siRNA           920         UUUCCUCGAAGUGCCAGUUUUCGU         110         sense         31535         KRAS2:999U21 siRNA           920         UUUCCUCGAAGUCGAUUGAUU         111         sense         31535         KRAS2:999U21 siRNA	OR	3892	UCACCUGUUUCCUGUAUGGAGGA	106	seuse	31436	KDR:3894U21 siRNA	ACCUGUUCCUGUAUGGAGTT
3302         UGACCUUGGAGCAUCUCGUCUCGUCUCGUCUCGUCGUCGAGCAUCCUCGUCGAGCAUCCUCGUCGAGCAUCCUCGUCGAGCAUCCUCGUCGAGGAUCCUCGUCUCGAGGAUCCUCGUCUCGAGGAUCCUCGUCUCGAGGAGGAGGAGGAGGGAUCAGGGAGGG	OR	3946	GACAACACAGCAGGAAUCAGUCA	107	seuse	31437	KDR:3948U21 siRNA	CAACACAGCAGGAAUCAGUTT
3852         UUUGAGCAUGGAAGAGGAUUCUG         102         antisense         31439         KDR:3872L21 siRNA           3892         UCACCUGUUUCCUGUAUGGAGGA         106         antisense         31440         KDR:3912L21 siRNA           3946         GACAACACAGCAGGAAUCAGUCA         107         antisense         31441         KDR:396L21 siRNA           625         ACAAGACAGGGUGUUGAUGAUGC         108         sense         31533         KRAS2:625U21 siRNA           920         UUUCCUCGAAGUGCCAGUAUUCC         108         sense         31533         KRAS2:625U21 siRNA           920         UUUCCUCGAAGUGCCAGUAUUCC         109         sense         31534         KRAS2:920U21 siRNA           920         UUUCCUCGAAGUGCCAGUUUUCGU         110         sense         31534         KRAS2:999U21 siRNA           999         AUUUCUGGCGUUUUUUGGGUUUUUGGU         110         sense         31535         KRAS2:999U21 siRNA           1013         GUUUUUGGUGCAUGCAGUUGAUU         111         sense         31536         KRAS2:1013U21 siRNA           1013         GUUUUUGGUGCAUGCAGUUGAUU         111         sense         31536         KRAS2:643L21 siRNA           643         ACAAGACAGGGUGUUGAUGAUGC         108         antisense         31537         KRAS2:643L21 siRNA <td>KDR (DR</td> <td>3302</td> <td></td> <td>105</td> <td>antisense</td> <td>31438</td> <td>KDR:3322L21 siRNA (3304C)</td> <td>AGAUGAGAUGCUCCAAGGUTT</td>	KDR (DR	3302		105	antisense	31438	KDR:3322L21 siRNA (3304C)	AGAUGAGAUGCUCCAAGGUTT
3892         UCACCUGUUUCCUGUAUGGAGGA         106         antisense         31440         KDR:3912L21 siRNA           3946         GACAACACAGCAGCAGCAGUCAUCAGUCA         107         antisense         31441         KDR:3966L21 siRNA           625         ACAAGACAGGGUGUUGAUGAUGC         108         sense         31533         KRAS2:625U21 siRNA           920         UUUCCUCGAAGUGCCAGUAUUCC         109         sense         31534         KRAS2:625U21 siRNA           920         UUUCCUCGAAGUGCCAGUAUUCC         109         sense         31534         KRAS2:625U21 siRNA           920         UUUCCUCGAAGUGCCAGUAUUCC         109         sense         31534         KRAS2:920U21 siRNA           999         AUUUCCUCGAAGUGCCAGUUUUUGGU         110         sense         31535         KRAS2:999U21 siRNA           1013         GUUUUUGGUGCAUGCAGUUGAUU         111         sense         31535         KRAS2:013U21 siRNA           1013         GUUUUUGGUGCAUGCAGUUGAUUGAUUG         111         sense         31536         KRAS2:643L21 siRNA           643         ACAAGACAGGGGUGUUGAUGAUGC         108         antisense         31537         KRAS2:643L21 siRNA	COR	3852	UUUGAGCAUGGAAGAGGAUUCUG	102	antisense	31439	KDR:3872L21 siRNA (3854C)	GAAUCCUCUUCCAUGCUCATT
3946         GACAACACAGCAGCAGUCAGUCA         107         antisense         31441         KDR:3966L21 siRNA           625         ACAAGACAGGGUGUUGAUGAUGC         108         sense         31533         KRAS2:625U21 siRNA           920         UUUCCUCGAAGUGCCAGUAUUCC         109         sense         31534         KRAS2:625U21 siRNA           920         UUUCCUCGAAGUGCCAGUAUUCC         109         sense         31534         KRAS2:920U21 siRNA           930         AUUUCCUCGAAGUGCCAGUAUUCGU         110         sense         31534         KRAS2:920U21 siRNA           999         AUUUCUGUCUUGGGGUUUUUGGU         110         sense         31535         KRAS2:999U21 siRNA           1013         GUUUUUGGUGCAUGCAGUUGAUU         111         sense         31535         KRAS2:1013U21 siRNA           1013         GUUUUUGGUGCAUGCAGUUGAUU         111         sense         31536         KRAS2:1013U21 siRNA           643         ACAAGACAGGGUGUUGAUGAUGC         108         antisense         31537         KRAS2:643L21 siRNA           643         ACAAGACAGGGUGUUGAUGAUGC         108         antisense         31537         KRAS2:643L21 siRNA	KDR	3892	UCACCUG	106		31440	KDR:3912L21 siRNA (3894C)	CUCCAUACAGGAAACAGGUTT
625         ACAAGACAGGGUGUUGAUGAUGC         108         sense         31533         KRAS2:625U21 siRNA         ACAAGACAGGGUGUUGAUGAUGC         108         sense         31533         KRAS2:625U21 siRNA         ACAAGACAGGGUGUUGAUGCC         109         sense         31534         KRAS2:920U21 siRNA         ACAAGACGGGUGUUGAUGCC         109         sense         31534         KRAS2:920U21 siRNA         URAS2:990U21 siRNA	KDR	3946	<u> </u>	107		31441	KDR:3966L21 siRNA (3948C)	ACUGAÜUCCUGCUGUGUUGT
625         ACAAGACAGGGUGUUGAUGAUGC         108         sense         31533         KRAS2:625U21 siRNA           920         UUUCCUCGAAGUGCCAGUAUUCC         109         sense         31534         KRAS2:920U21 siRNA           920         UUUCCUCGAAGUGCCAGUAUUCC         109         sense         31534         KRAS2:920U21 siRNA           999         AUUUCUGUCUUGGGGUUUUUGGU         110         sense         31535         KRAS2:999U21 siRNA         U           1013         GUUUUUGGUGCAUGCAGUUGAUU         111         sense         31536         KRAS2:1013U21 siRNA         U           1013         GUUUUUGGUGCAUGCAGUGAUU         111         sense         31536         KRAS2:1013U21 siRNA         U           643         ACAAGACAGGGUGUUGAUGAUGC         108         antisense         31537         KRAS2:643L21 siRNA           643         ACAAGACAGGGUGUUGAUGAUGC         108         antisense         31537         KRAS2:643L21 siRNA	RAS2	625	ACAAGACAGGGUGUUGAUGAUGC	108	seuse	31533	KRAS2:625U21 siRNA	AAGACAGGGUGUUGAUGAUTT
920         UUUCCUCGAAGUGCCAGUAUUCC         109         sense         31534         KRAS2:920U21 siRNA           920         UUUCCUCGAAGUGCCAGUAUUCC         109         sense         31534         KRAS2:920U21 siRNA         1           999         AUUUCUGUCUUGGGGUUUUUGGU         110         sense         31535         KRAS2:999U21 siRNA         L           999         AUUUCUGUCUUGGGUUUUGGU         111         sense         31535         KRAS2:999U21 siRNA         L           1013         GUUUUUGGUGCAUGCAGUUGAUU         111         sense         31536         KRAS2:1013U21 siRNA         L           643         ACAAGACAGGGUGUUGAUGAUGC         108         antisense         31537         KRAS2:643L21 siRNA         L           643         ACAAGACAGGGUGUUGAUGAUGC         108         antisense         31537         KRAS2:643L21 siRNA         L	RAS2	625	ACAAGACAGGGUGUUGAUGAUGC	108	sense	31533	KRAS2:625U21 siRNA	AAGACAGGGUGUUGAUGAUTT
920         UUUCCUCGAAGUGCCAGUAUUCC         109         sense         31534         KRAS2:920U21 siRNA           999         AUUUCUGUCUUGGGGUUUUUUGGU         110         sense         31535         KRAS2:999U21 siRNA         U           1013         AUUUCUGUCUUGGGGUUUUUGGU         111         sense         31535         KRAS2:999U21 siRNA         U           1013         GUUUUUUGGUGCAUGCAUGAUU         111         sense         31536         KRAS2:1013U21 siRNA         U           643         ACAAGACAGGGUGUUGAUGAUGC         108         antisense         31537         KRAS2:643L21 siRNA         U           643         ACAAGACAGGGUGUUGAUGAUGC         108         antisense         31537         KRAS2:643L21 siRNA	RAS2	ı	UNUCCUCGAAGUGCCAGUAUUCC	109	esues	31534	KRAS2:920U21 siRNA	UCCUCGAAGUGCCAGUAUUTI
999         AUUUCUGUCUUGGGGUUUUUGGU         110         sense         31535         KRAS2:999U21 siRNA         U           999         AUUUCUGUCUUGGGGUUUUUUGGU         110         sense         31535         KRAS2:999U21 siRNA         U           1013         GUUUUUGGUGCAUGCAUGAUU         111         sense         31536         KRAS2:1013U21 siRNA         U           1013         GUUUUUGGUGCAUGCAUGAUGAU         111         sense         31536         KRAS2:1013U21 siRNA         U           643         ACAAGACAGGGUGUUGAUGAUGC         108         antisense         31537         KRAS2:643L21 siRNA           643         ACAAGACAGGGUGUUGAUGAUGC         108         antisense         31537         KRAS2:643L21 siRNA	RAS2		UNUCCUCGAAGUGCCAGUAUUCC	109	seuse	31534	KRAS2:920U21 siRNA	UCCUCGAAGUGCCAGUAUUTT
999         AUUUCUGUCUUGGGGUUUUUGGU         110         sense         31535         KRAS2:999U21 siRNA         U           1013         GUUUUUGGUGCAUGCAGUUGAUU         111         sense         31536         KRAS2:1013U21 siRNA         U           1013         GUUUUUGGUGCAUGCAGUUGAUGAUGC         108         antisense         31536         KRAS2:643L21 siRNA         U           643         ACAAGACAGGGUGUUGAUGAUGC         108         antisense         31537         KRAS2:643L21 siRNA	RAS2	<u> </u>	AUUUCUGUCUUGGGGUUUUUGGU	110	esues	31535	KRAS2:999U21 siRNA	uncuencuneeeeuuuuuet
1013         GUUUUUGGUGCAUGCAGUUGAUU         111         sense         31536         KRAS2:1013U21 siRNA         1           1013         GUUUUUGGUGCAUGCAUGCAGUUGAUU         111         sense         31536         KRAS2:1013U21 siRNA         0           643         ACAAGACAGGGUGUUGAUGAUGAUGC         108         antisense         31537         KRAS2:643L21 siRNA           643         ACAAGACAGGGUGUUGAUGAUGAUGC         108         antisense         31537         KRAS2:643L21 siRNA	RAS2	l	AUUUCUGUCUUGGGGUUUUUGGU	110	esues	31535	KRAS2:999U21 siRNA	UNCUGUCUUGGGGUUUUUGT
1013         GUUUUUGGUGCAUGCAGUUGAUU         111         sense         31536         KRAS2:1013U21 siRNA         1           643         ACAAGACAGGGUGUUGAUGAUGC         108         antisense         31537         KRAS2:643L21 siRNA           643         ACAAGACAGGGUGUUGAUGAUGC         108         antisense         31537         KRAS2:643L21 siRNA	RAS2	1013	-	111	sense	31536	KRAS2:1013U21 siRNA	UUUUGGUGCAUGCAGUUGAT
643         ACAAGACAGGGUGUUGAUGAUGC         108         antisense         31537         KRAS2:643L21 siRNA           643         ACAAGACAGGGUGUUGAUGAUGC         108         antisense         31537         KRAS2:643L21 siRNA	RAS2		GUUUUUGGUGCAUGCAGUUGAUU	111	sense	31536	KRAS2:1013U21 siRNA	UUUUGGUGCAUGCAGUUGAT
643 ACAAGACAGGGUGUUGAUGAUGC 108 antisense 31537 KRAS2:643L21 siRNA	RAS2	643	ACAAGACAGGGUGUUGAUGAUGC	108	antisense	31537	KRAS2:643L21 siRNA (625C)	AUCAUCAACACCCUGUCUUII
	RAS2		ACAAGACAGGGUGUUGAUGAUGC	108	antisense	31537	KRAS2:643L21 siRNA	AUCAUCAACACCCUGUCUUTT

(+01 NO+)

·	-	ч
١	ſ	7
	•	٠.
۳	-	٦

089	681	682	683	684	685	3	989		687	989	689	900	069	691	692		693	804		695	909	080	/69 69/	869	669	9	79	200	70/	703	_
AAGAGCUCCAGCUUAGAGUTT	CAGCUUGGAACACCAUGUCTT	IIICCCAGCUGACUCAGAACTT	ALIGICAACAGAUCCGACUUTT	HIGGOODICHGAIACAGACATT	TEGINORYONING	GACAUGGUGUUCCAAGCUGII	GUICUGAGUCAGCUGGGAATT		AAGUCGGAUCUGUUGACAUTT	UGUCUGUAUCAGAGGCCAATT	F C. CACCACCALIGAGGATT B		B GGucuGuuAuuGccAAGcATT B	B cuGcAGGAGucuucAAAAGTT B	B TOOK ON THE B	B GugcuAccAAcAcAcAcaa B	uccucAucAuccucGucAcTsT	H-	uGcuuGGcAAuAAcAGAcc181	GuiningAAGAGICCUGCAGTST	+	GGuucuGuGuuGGuAGcACISI	GUGACGAGGAUGAUGAGGATT	GGUCUGUUAUUGCCAAGCATT	CUGCAGGAGUCUUCAAAAGTT	GUGCUACCAACACAGAACCTT	LICCUCAUCAUCCUCGUCACTT		UGCUUGGCAAUAACAGACCII	CUUUUGAAGACUCCUGCAGTT	
(2884C) 31593 MAPK14:3574L21 siRNA	(3556C) 04547 MADKR:7351124 siRNA	WAPAGE SOCE SHARE	31518 MAPK8:855021 SININ	31519 MAPK8:1220021 SININA	31520 MAPK8:1244UZ1 SIKINA	antisense 31521 MAPK8:753L21 siRNA	(/35C)	31522 MAPK8:873LZ1 SIKINA (855C)	31523 MAPK8:1244L21 siRNA	31524 MAPK8:1262L21 siRNA	(1244C)	30977 MYB:148U21 siRNA stab04	30978 MYB:457U21 siRNA	30979 MYB:708U21 siRNA	stab04	30980 MYB:1053U21 siRNA	20081 MVR:1661 21 siRNA	(148C) stab05	30982 MYB:475L21 siRNA	(457C) stab05	30983 MYB:/26LZ1 SIRINA (708C) stab05	30984 MYB:1071L21 siRNA	34025 MYB:148U21 siRNA	31026 MVR-457U21 siRNA	31020 MVB-708121 SIRNA	31021 WILLIAGOZI SII. S. 34028 MVB:10531 121 SIBNA	MIT B. 103302 1 31 KK	31101 MYB: 100L2 1 SINNA (148C)	31102 MYB:475L21 siRNA	(457C)	antisense (31103 MYB:/zocz (31104 MYB:/z
31593	04547	31517	31518	31519	31520	31521			31523			30977	30978	30079		30980	2000						3402	3402	2102	2102					9 3110
antisense		seuse	sense	seuse	sense	antisense		antisense	antisense	antisense		seuse	sense	0000	20120	seuse		aunzenze	antisense		antisense	antisense	00000	201120	Selise	Selise	seuse	antisense	antisense		antisens
119	9	120	121	122	123	120		121	122	123	2	57	57	57	õ	22	1	<u> </u>	57		29	22	707	124	2 3	120	127	124	125		126
GGACUCUAAGCUGGAGCUCUUGG		AACAGCUUGGAACACCAUGUCCU	UNUNCCCAGCUGACUCAGAACAC	CAAUGUCAACAGAUCCGACUUUG	CHILIPECCHICHGALIACAGACAGC	AACAGCUUGGAACACCAUGUCCU		UNUUCCCAGCUGACUCAGAACAC	CAALIGLICAACAGAUCCGACUUUG	1		AACAACCACAAAAUACAACAAGA	AACAACCACAAAAUACAACAAGA		AACAACCACAAAAUACAACAAGA	AACAACCACAAAAUACAACAAGA		AACAACCACAAAAUACAACAAGA	AACAACCACAAAAJACAACAAGA		AACAACCACAAAAUACAACAAGA	AACAACCACAAAAUACAACAAGA	-			AUCUGCAGGAGUCUUCAAA	ـــ	┼	I II GELICI GUI BUI GCCAAGCACU		AUCUGCAGGAGUCUUCAAAAGCC
3574		733	853	1224	1040	733	 }	853	1224	1771	1242	146	455	3	902	1051		146	AFF	5	902	1051		146	455	902	1051	146	707		706
4 MAPK1		MAPK8			0/10/4	MAPK8	2	MAPK8	MADKR		MAPK8	MYB	MVB	2	MYB	MYB	1	MYB	QVV	<u> </u>	МУВ	MYB		MYB	MYB	MYB	MYB	MYB		<u> </u>	MYB

PCT/US03/05028

4	
9	
-	

704	705	902	707	802	709	710	711	710	71.2	717	745	716	2 !	7117	718	719	700	02)	80/	712	721	722	100	67)	47)
GGUUCUGUGUUGGUAGCACTT	B AGAGGGucAAGuuGGAcAGTT B	B GCAGAGGAGCAAAAGCUCATT B	B cGGAAcucuuGuGcGuAAGTT B	B AAccuuGGcuGAGucuuGATT B	cuGuccAAcuuGAccucuTsT	Talladiadiadiamon	Total Control of the	cuuAcGcAcAAGAGuuccGISI	ucAAGAcucAGccAAGGuuls1	AGAGGGUCAAGUUGGACAGII	GCAGAGGAGCAAAAGCUCAII	CGGAACUCUUGUGCGUAAGII	AACCUUGGCUGAGUCUUGALI	CUGUCCAACUUGACCCUCUTT	UGAGCUUUUGCUCCUCUGCTT	CHITACGCACAAGAGUUCCGTT		UCAAGACUCAGCCAAGGUUII	B AAccuuGGcuGAGucuuGATT B	ucAAGAcucAGccAAGGuuTsT	B AAccuuGGcuGAGucuuGATT B	TeTOGAACCAACAACAA	ווכאאופאפונאפונאפונאפונאפונאפונאפונאפונאפונ	B AGuucuGAGucGGuuccAA11 B	uuGGAAccGAcucAGAAcuTsT
antisense 31104 MYB:1071L21 siRNA	30825 MYC:1526U21 siRNA	30826 MYC:1780U21 siRNA	30827 MYC:1861U21 siRNA	30828 MYC:1971U21 siRNA	30829 MVC:1544  21 siRNA	(1526C) stab05	30830 MYC:1798L21 siRNA (1780C) stab05	30831 MYC:1879L21 siRNA (1861C) stab05	30832 MYC:1989L21 siRNA (1971C) stab05	30993 MYC:1526U21 siRNA	30994 MYC:1780U21 siRNA	30995 MYC:1861U21 siRNA	20096 MYC:1971U21 siRNA	31069 MYC:1544L21 siRNA	(1526C) antisense 31070 MYC:1798L21 siRNA	(1780C)	31071 MYC:1879L21 SIKNA (1861C)	31072 MYC:1989L21 siRNA	31377 MYC:1971U21 siRNA	31380 MYC:1989L21 siRNA	(1971C) stab05	\$1365 WITC. 137 102.1 SHEET	31386 MYC:1989L21 siRNA (1971C) stab11	31389 MYC:1971U21 siRNA inv	antisense 31392 MYC:1989L21 siRNA (1971C) inv stab05
antisense 3	sense 3	sense 3	sense 3	sense		allisalisa	antisense	antisense	antisense	sense	+	$\top$	+	antisense	antisense		antisense	antisense	seuse	antisense		seuse	antisense	seuse	antisense
127	128	129	130	131	000	971	129	130	131	128	120	3 8	3 5	128	129		130	131	131	131	-	131	131	131	131
AGGUGCUACCAACACAGAACCAC	CAAGAGGGUCAAGUUGGACAGUG	AAGCAGAGGAGCAAAAGCUCAUU	INCERACTICI II GI IGCGI I AAGGA	ACAACCIIIGGCIIGAGUCUUGAGA		CAAGAGGGUCAAGUUGGACAGUG	AAGCAGAGGAGCAAAAGCUCAUU	UACGGAACUCUUGUGCGUAAGGA	ACAACCUUGGCUGAGUCUUGAGA	STATE OF THE PROPERTY OF THE P	CAAGAGGGCCAAGCGGGACAGGC	AAGCAGAGCAGCAAAAGCACACACACACACACACACACA	UACGGAACUCUUGUGCGCAAAGAA	ACAACCUUGGCUGAGUCUUGAGA CAAGAGGGUCAAGUUGGACAGUG			UACGGAACUCUUGUGCGUAAGGA	ACAACCUUGGCUGAGUCUUGAGA		1		ACAACCUUGGCUGAGUCUUGAGA	ACAACCUUGGCUGAGUCUUGAGA	ACAACCUUGGCUGAGUCUUGAGA	ACAACCUUGGCUGAGUCUUGAGA
1051	-+-					1524	1778	1859	1969		1524	17.78	1859	1524	3	1778	1859	1969	1969			1969	1969	1969	
МУВ	MYC	CAM	2 2		2	MYC	MYC	MYC	MYC		MYC	MYC	MYC	MYC	2	MYC	MYC	MYC	MYC		ک آج 	MYC	MYC	MYC	MYC

725	726	727	728	729	730	731	732	733	734	735	736	737	738	739	740	741	742	743	744	745
B AGuucuGAGucGGuuccAATT B	uuGGAAccGAcucAGAAcuTsT	B GuucAGuGucucuccAAAATT B	B uuuGcAGAuAGccuuGAGcTT B	B uccuGcuGcuuucAuuGAcTT B	B GAcuGccAuGuGuucAucATT B	uuuuGGAGAGAcAcuGAAcTsT	GcucAAGGcuAucuGcAAATsT	GucAAuGAAAGcAGcAGGATsT	uGAuGAAcAcAuGGcAGucTsT	CUGCAGUACCUCUACCUGCTT	CGUCUCCUACUGCACCAGATT	UGGAGCCUGGAAGACCAGCTT	GUGACUCAGAAGGCUCAGGTT	GCAGGUAGAGGUACUGCAGTT	UCUGGUGCAGUAGGAGACGTT	GCUGGUCUUCCAGGCUCCATT	CCUGAGCCUUCUGAGUCACTT	B uGcAcGuAuAuGccGAGAuTT B	B ccAuAuuGGAGAuGcuGuuTT B	B AuuGcGGAuAuGGGAcAcuTT B
31395 MYC:1971U21 siRNA inv stab07	38 MYC:1989L21 siRNA (1971C) inv stab11	30833 Nogo:1043U21 siRNA stab04	30834 Nogo:1407U21 siRNA stab04	30835 Nogo:3211U21 siRNA stab04	30836 Nogo:3883U21 siRNA stab04		30838 Nogo:1425L21 siRNA (1407C) stab05	39 Nogo:3229L21 siRNA (3211C) stab05	t0 Nogo:3901L21 siRNA (3883C) stab05	57 NogoR:512U21 siRNA	31058 NogoR:662U21 siRNA	31059 NogoR:1086U21 siRNA	31060 NogoR:1371U21 siRNA	31133 NogoR:530L21 siRNA (512C)	31134 NogoR:680L21 siRNA (662C)	31135 NogoR:1104L21 siRNA (1086C)	36 NogoR:1389L21 siRNA (1371C)	11 PCNA:550U21 siRNA stab04	12 PCNA:574U21 siRNA stab04	44 PCNA:839U21 siRNA stab04
	se 31398		1	<b></b>		se 30837	<del> </del>	se 30839	se 30840	31057		1	t -				lse 31136	a 30841	9 30842	e 30844
seuse	antisense	seuse	seuse	sense	seuse	antisense	antisense	antisense	antisense	seuse	esues	seuse	seuse	antisense	antisense	antisense	antisense	seuse	seuse	seuse
131	131	132	133	134	135	132	133	134	135	136	137	138	139	136	137	138	139	140	141	142
1969 ACAACCUUGGCUGAGUCUUGAGA	ACAACCUUGGCUGAGUCUUGAGA	UCGUUCAGUGUCUCCCAAAAGC	GUUUUGCAGAUAGCCUUGAGCAA	AUUCCUGCUGCUUCAUUGACAG	UUGACUGCCAUGUGUUCAUCAUC	UCGUUCAGUGUCUCCCAAAAGC	GUUUUGCAGAUAGCCUUGAGCAA	AUUCCUGCUGCUUCAUUGACAG	UUGACUGCCAUGUGUUCAUCAUC	CCCUGCAGUACCUCUACCUGCAG	ACCGUCUCCUACUGCACCAGAAC	NOGO 1084 ACUGGAGCCUGGAAGACCAGCUU	1369 UGGUGACUCAGAAGGCUCAGGUG	cccuecaeuaccucuaccuecae	ACCGUCUCCUACUGCACCAGAAC	ACUGGAGCCUGGAAGACCAGCUU	UGGUGACUCAGAAGGCUCAGGUG	UUUGCACGUAUAUGCCGAGAUCU	AGCCAUAUUGGAGAUGCUGUUGU	AAAUUGCGGAUAUGGGACACUUA
	1969	1043	1407	3211	3883	1061	1425	3229	3901	510	099	1084	1369	510	099	1084	1369	548	572	837
MYC	MYC	Nogo	Nogo	Nogo	Nogo	Nogo	Nogo	Nogo	Nogo	NOGO R	NOGO R	NOGO R	NOGO R	NOGO R	NOGO R	NOGO R	NOGO R	PCNA	PCNA	PCNA

`	0
Ĺ	Ó
÷	Ξ
•	

746	747	748	749	750	751	752	753	754	101	3	96/	757	758	759	760	Ş	ē	761	762	762	762	762	763	3 6	20
AucucGGcAuAuAcGuGcATsT	AAcAGcAucuccAAuAuGGTsT	AGuGucccAuAuccGcAAuTsT	LIGCACGUAUAUGCCGAGAUTT	CCALIAUUGGAGAUGCUGUUTT	AAAGCCACUCCACUCUUTT	AUUGCGGAUAUGGGACACUTT	AUCUCGGCAUAUACGUGCATT	TTOOLINILANDOLINITADOLINI	AACAGCAUCUCCAAOAOGGTT	AAGAGAGUGGAGUGGCUUUTT	AGUGUCCCAUAUCCGCAAUTT	B AAAGccAcuccAcucuuTT B	AAGAGAGuGGAGuGGcuuuTsT	B uucucucAccucAccGAAATT B	HILLINGGGUGAGGUGAGAGATST		B cAGGAccuccAcAuGAuAG11 B	B cAGGAccuccAcAuGAuAGTT B	B AGAuuuGAccuuccuGAcATT B	B AGAuuuGAccuuccuGAcATT B	B AGAuuuGAccuuccuGAcATT B	B AGAuuuGAccuuccuGAcATT B	B TTOOKSAACCCX. C.K.	B uGAGuAGCUGGAUUACAGGTT B	B uGAGuAGcuGGAuuAcAGG11 B
antisense 30845 PCNA;568L21 siRNA	30846 PCNA:592L21 siRNA		(839C) stabus	31033 PCINA:3300Z1 SINIA	31034 PCNA:3740Z1 SINNA	31035 PCIMA: 701 021 311 NA	Sense 31030 FONA:3502 Single	(550C)	31110 PCNA:592L21 siRNA (674C)			31310 PCNA:767U21 siRNA	31311 PCNA:785L21 siRNA	31322 PCNA:767U21 siRNA inv	stab04	(767C) inv stab05	30969 PKR:533U21 siRNA	30969 PKR:533U21 siRNA	70 PKR:1171U21 siRNA	70 PKR:1171U21 siRNA	30970 PKR:1171U21 siRNA	30970 PKR:1171U21 siRNA	stab04	71 PKR:2430U21 siRNA stab04	30971 PKR:2430U21 sIRNA stab04
3084		30848	3	3103	3103	3103	010	01.TS		31111	3111	3131		3132		31323	3606	309(	30970	30970	309			30971	
antisense	antisense	antisense		sense	seuse	seuse	asuas	antisense	antisense	antisense	antisense 31112	sense	antisense	sense		antisense	seuse	seuse	seuse	seuse	sense	asuas	5	seuse	seuse
140	141	142	1	64	141	143	142	140	141	143	142	143	143	143		143	144	144	22	22	22	57	5	22	22
UNUGCACGUAUAUGCCGAGAUCU	AGCCAUAUUGGAGAUGCUGUUGU	AAAUUGCGGAUAUGGGACACUUA		UNUGCACGUAUAUGCCGAGAUCU	AGCCAUAUUGGAGAUGCUGUUGU	CAAAAGCCACUCCACUCUCA	AAAUUGCGGAUAUGGGACACUUA	UUUGCACGUAUAUGCCGAGAUCU	AGCCAUAUUGGAGAUGCUGUUGU	CAAAAGCCACUCCACUCUCU	AAAUUGCGGAUAUGGGACACUUA	CAAAAGCCACUCCACUCUCA	CAAAAGCCACUCCACUCUCA	CAAAAGCCACIICCACIICUCUCA		CAAAAGCCACUCCACUCUCUCA	UUCAGGACCUCCACAUGAUAGGA	UUCAGGACCUCCACAUGAUAGGA	AACAACCACAAAAUACAACAAGA		$\dashv$		AACAACCACAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA	AACAACCACAAAAUACAACAAGA	AACAACCACAAAAUACAACAAGA
548	572	837		548	572	765	837	548	572	765	837	765	765	785	3	765	533	533	1171	1171	1171		1171	2430	2430
PCNA	PCNA	PCNA	· : :	PCNA	PCNA	PCNA	PCNA	PCNA	PCNA	PCNA	PCNA	PCNA	PCNA	V 1400	5	PCNA	PKR	PKR	PKR	PKR	0/10		PKR	PKR	PKR

ù	2	
	=	

764	764	765	765	992	992	767	767	768	768	769	770	771	772	773	774	775	776	777	778	779	780	781
B GGucucAAAcuccuGAccuTT B	B GGucucAAcuccuGAccuTT B	cuAucAuGuGGAGGuccuGTsT	cuAucAuGuGGAGGuccuGTsT	uGucAGGAAGGucAAAucuTsT	uGucAGGAAGGucAAAucuTsT	ccuGuAAuccAGcuAcucATsT	ccuGuAAuccAGcuAcucATsT	AGGucAGGAGuuuGAGAccTsT	AGGucAGGAGuuuGAGAccTsT	B AAAGGcuGAGGuuGcuGAuTT B	B AAAcAAccuuccAAcAAcTT B	B AAGGAcuGAuGAccAAAcATT B	AucAGcAAccucAGccuuuTsT	GGuuGuuGGAAGGuuGuuTsT	uGuuuGGucAucAGuccuuTsT	AAAGGCUGAGGUUGCUGAUTT	AAACAACCUUCCAACAACCTT	GGAUGUGGUGAUUCAGGAUTT	AAGGACUGAUGACCAAACATT	AUCAGCAACCUCAGCCUUUTT	GGUUGUUGGAAGGUUGUUUTT	AUCCUGAAUCACCACAUCCTT
30972 PKR:2518U21 siRNA stab04	30972 PKR:2518U21 siRNA stab04	30973 PKR:551L21 siRNA (533C) stab05	30973 PKR:551L21 siRNA (533C) stab05	30974 PKR:1189L21 siRNA (1171C) stab05		30975 PKR:2448L21 siRNA (2430C) stab05	30975 PKR:2448L21 siRNA (2430C) stab05	76 PKR:2536L21 siRNA (2518C) stab05	76 PKR:2536L21 siRNA (2518C) stab05		14 PRKCA:1000U21 siRNA stab04	30716 PRKCA:1736U21 siRNA stab04	17 PRKCA:537L21 siRNA (519C) stab05		20 PRKCA:1754L21 siRNA (1736C) stab05	30989 PRKCA:519U21 siRNA	30990 PRKCA:1000U21 siRNA	30991 PRKCA:1143U21 siRNA	92 PRKCA:1736U21 siRNA	65 PRKCA:537L21 siRNA (519C)	31066 PRKCA:1018L21 siRNA (1000C)	
3097	3097	+		+	30974		<del>                                     </del>	30976	30976	30713	30714	307	9 30717	9 30718		T	308	308	30992	a 31065	1	e 31067
sense	sense	antisense	antisense	antisense	antisense	antisense	antisense	antisense	antisense	seuse	sense	sense	antisense	antisense	antisense	sense	seuse	sense	seuse	antisense	antisense	antisense
57	57	22	22	57	22	57	22	22	57	145	146	147	145	146	147	145	146	148	147	145	146	148
AACAACCACAAAAUACAACAAGA	AACAACCACAAAAUACAACAAGA	AAGAACCACAAAAUACAACAAGA	AACAACCACAAAAUACAACAAGA	AACAACCACAAAAUACAACAAGA	AACAACCACAAAAUACAACAAGA	AACAACCACAAAAUACAACAAGA	AACAACCACAAAAUACAACAAGA	AACAACCACAAAAUACAACAAGA	AACAACCACAAAAUACAACAAGA	CUAAAGGCUGAGGUUGCUGAUGA	GGAAACAACCUUCCAACAACCUU	CAAAGGACUGAUGACCAAACACC	CUAAAGGCUGAGGUUGCUGAUGA	GGAAACAACCUUCCAACAACCUU	CAAAGGACUGAUGACCAAACACC	CUAAAGGCUGAGGUUGCUGAUGA	GGAAACAACCUUCCAACAACCUU	4		CUAAAGGCUGAGGUUGCUGAUGA	GGAAACAACCUUCCAACAACCUU	AAGGAUGUGGUGAUUCAGGAUGA
2518	2518	551	551	1189	1189	2448	2448	2536	2536	517	866	1734	517	866	1734	517	866	1141	1734	517	866	1141
PKR	PKR	PKR	PKR	RX TX	PKR	PKR	PKR	PKR	PKR	PRKCA	PRKCA	PRKCA	PRKCA	PRKCA	PRKCA	PRKCA	PRKCA	PRKCA	PRKCA	PRKCA	PRKCA	PRKCA

٠		•
١	ſ	1
•	٠.	•
•	-	-

782	783	784	785	786	787	788	789	200	08.	791	792	793	794	795	796	101	/8/	798	199	800	801	3	802
UGUUUGGUCAUCAGUCCUUTT	B GGAuGuGGuGAuucAGGAuTT B	AuccuGAAucAccAcAuccTsT	B GGAuGuGGuGAuucAGGAuTT B	AuccuGAAucAccAcAuccTsT	B uAGGAcuuAGuGGuGuAGGTT B	CCUACACCACUAAGuccuATsT	B TION COLUMNIA GIRGING IN A GIGTT B	E LAGGACUUA CUA CUA CUA CUA CUA CUA CUA CUA CU	ccuAcAccAcuAAGuccuAISI	CUCGUUUCUCUUGGACAAGTT	GGUGAGCUACAAACACAUGTT	AGUGGAAGACUGGCUGAGCTT	CCUCUAGCCUGUUGUTT	CILICIAAGAGAAACGAGTT		CAUGUGUUGGAGCACACA	GCUCAGCCAGUCUUCCACUTI	ACAACAACAGGCUAGAGGTT	B uccGAcAuGAAGccAGuGATT B	B CHICALIGGACAAGAGGAAAGTT B		B GuGuGGAuAAGGcuuAGcuii B	ucAcuGGcuucAuGucGGA181
antisense 31068 PRKCA:1754L21 siRNA	(1736C) 31376 PRKCA:1143U21 siRNA	31379 PRKCA:1161L21 siRNA	(1143C) stab05 31382 PRKCA:1143U21 siRNA	31385 PRKCA:1161L21 siRNA	31388 PRKCA:1143U21 siRNA	inv stab04	31391 PRKCA:1161L21 SIRNA (1143C) inv stab05	31394 PRKCA:1143U21 siRNA inv stab07	31397 PRKCA:1161L21 siRNA	31557 PTP4A3:205U21 siRNA	31558 PTP4A3:367U21 siRNA	31559 PTP4A3:574U21 siRNA	31560 PTP4A3:1168U21 siRNA		31561 PTP4A3:223L21 SIKNA (205C)	31562 PTP4A3:385L21 siRNA	31563 PTP4A3:592L21 siRNA	31564 PTP4A3:1186L21 siRNA	30865 PTPN1:242U21 siRNA	stab04	30867 PTPN1:874U21 SIKINA stab04	30868 PTPN1:3037U21 siRNA stab04	30869 PTPN1:260L21 siRNA (242C) stab05
Infisense 31	sense 31	antisense 37		antisense 3	sense 3		antisense 3	sense 3	antisense 3	sense 3	sense 3	sense 3			antisense	antisense	antisense	antisense	Sense		seuse	seuse	antisense
147   8	148	-	148	148	148		148	148	148	149	150	151	153	70	149	150	151	152	153	3	154	155	153
PBVCA 1734   CAAAGGACIIGALIGACCAAACACC	+	AAGGALIGIIGGIIGAUUCAGGAUGA	AAGGALIGILGGLIGAUUCAGGAUGA	GAUGA			PRKCA 1141 AAGGAUGUGGUGAUUCAGGAUGA	PRKCA 1141 AAGGAUGUGGUGAUUCAGGAUGA						CUCCUCUAGCCUGUUGUUGUGG	AUCUCGUUUCUCUUGGACAAGCA	GAGGUGAGCUACAAACACAUGCG				UAUCCGACAUGAAGCCAGGGACG	UGCUGAUGGACAAGAGAGAC	5 AGGUGUGGAUAAGGCUUAGGUGC	UAUCCGACAUGAAGCCAGUGACU
173/	4444	1444	1411	1 1 1		114.	1141	1141	1141	205			1 2/4	1168	, 223	385			5	1 240	1 872	1 3035	1 240
VJAGO	101 101 101 101 101 101 101 101 101 101	7 Y	PRINCE	TANCA CONGO	אאר	PKKCA 1141	PRKCA	PRKCA	PRKCA	OTDAA	3 47	3 47	Р 1 Р 4 А 3	PTP4A	PTP4A	PTP4A	3 DTD4A	3 4	3	PTPN1	PTPN1	PTPN1	PTPN1

803	804	805	908	807	808	808	810		811	812	813	814	815	816	817	818	819	820	821	822	823	824	825	826	827	828	829
cunuccucuuGuccAucAGTsT	AccuAAGccuuAuccAcAcTsT	UCCGACAUGAAGCCAGUGATT	GUCCGAGAGUCAGGGUCACTT	CUGAUGGACAAGAGGAAAGTT	GUGUGGAUAAGGCUUAGGUTT	UCACUGGCUUCAUGUCGGATT	GUGACCCUGACUCUCGGACTT	***************************************	CUUUCCUCUUGUCCAUCAGTI	ACCUAAGCCUUAUCCACACTT	B GuccGAGAGucAGGGucAcTT B	GuGAcccuGAcucucGGAcTsT	B cAcuGGGAcuGAGAGccuGTT B	cAGGcucucAGucccAGuGTsT	AACACGGCAUGUGAACAUUTT	UCUACAACACCUGCAUGUTT	UCACAUCAACAACCGAGAUTT	AGGAAGCCAGGAAUACAGGTT	AAUGUUCACAUGCCGUGUUTT	ACAUGCAGGUGUUUGUAGATT	AUCUCGGUUGUUGAUGUGATT	CCUGUAUUCCUGGCUUCCUTT	GAGGAGCACAGAUACCACCTT	UGGCUUCUAUGAGGCUGAGTT	UGUGACAAGGUGCAGAAAGTT	CUCCAGCUUCUGGUACUCUTT	
antisense 30871 PTPN1:892L21 siRNA (874C) stab05	30872 PTPN1:3055L21 siRNA (3037C) stab05	31017 PTPN1:242U21 siRNA	31018 PTPN1:766U21 siRNA	31019 PTPN1:874U21 siRNA	31020 PTPN1:3037U21 siRNA	31093 PTPN1:260L21 siRNA	31094 PTPN1:784L21 siRNA	7 (766C)	31095 PTPN1:892L21 siRNA (874C)	31096 PTPN1:3055L21 siRNA (3037C)	31306 PTPN1:766U21 siRNA stab04	31307 PTPN1:784L21 siRNA (766C) stab05	31318 PTPN1:766U21 siRNA inv	31319 PTPN1:784L21 siRNA (766C) inv stab05	31549 RAF1:1326U21 siRNA	31550 RAF1:1415U21 siRNA	31551 RAF1:1776U21 siRNA	31552 RAF1:2854U21 siRNA	31553 RAF1:1344L21 siRNA (1326C)	31554 RAF1:1433L21 siRNA (1415C)	31555 RAF1:1794L21 siRNA (1776C)	31556 RAF1:2872L21 siRNA (2854C)	31029 ReIA:146U21 siRNA	31030 ReIA:290U21 siRNA	31031 ReIA:645U21 siRNA	31032 RelA:1957U21 siRNA	A140: 40 1404 A1 C 1044 A
antisense 3	antisense 3	seuse	1		1	e e	antisense		antisense	antisense	seuse	antisense	seuse	antisense	sense	$\top$	†	$\top$	antisense	antisense	antisense	antisense	seuse	1	1	T	
154	155	153	156	154	155	153	156		154	155	156	156	156	156	157	158	159	160	157	158	159	160	161	162	163	164	
UGCUGAUGGACAAGAGAAGAC	AGGUGUGGAUAAGGCUUAGGUGC	UAUCCGACAUGAAGCCAGUGACU	AAGUCCGAGAGUCAGGGUCACUC	UGCUGAUGGACAAGAGGAAAGAC	1 7		AAGUCCGAGAGUCAGGGUCACUC		UGCUGAUGGACAAGAGGAAAGAC	AGGUGUGGAUAAGGCUUAGGUGC	AAGUCCGAGAGUCAGGGUCACUC	AAGUCCGAGAGUCAGGGUCACUC	AAGUCCGAGAGUCAGGGUCACUC	AAGUCCGAGAGUCAGGGUCACUC	AAAACACGGCAHGHGAACAUHCU	CCHCHACAAACACCHGCAHGUCC			AAAACACGGCAUGUGAACAUUCU	CCUCUACAAACACCUGCAUGUCC	UCUCACAUCAACAACCGAGAUCA	CAAGGAAGCCAGGAAUACAGGUU	GAGAGGACACAGAUACCACCAA	GALIGGCIIICIJALIGAGGCUGAGCU	HGHGHGACAAGGUGCAGAAAGAG	_	4
872	3035	240	764	872			764		872	3035	764	764	764	764	1326	1415	1776	2854	1344	1433	1794	2872	144	288	643	1955	2
PTPN1	PTPN1	PTPN1	PTPN1	PTPN1	PTPN1	PTPN1	PTPN1	: : :	PTPN1	PTPN1	PTPN1	PTPN1	PTPN1	PTPN1	PAF1	RAF1	PAF1	RAF1	RAF1	RAF1	RAF1	RAF1	RFI A	\ <u>\</u>	REI A	RELA	

	830	831	832	833	834	835	836	837	838	839	840	841	842	843	0.44	044	845	846	847	848	φ 94	820	851	3
	CUCAGCCUCAUAGAAGCCATT	CUUUCUGCACCUUGUCACATT	AGAGUACCAGAAGCUGGAGTT	B cuccAGcuucuGGuAcucuTT B	AGAGuAccAGAAGcuGGAGTsT	B ucucAuGGucuucGAccucTT B	GAGGucGAAGAccAuGAGATsT	B uAuGcuGuGGuGcuuAAuGTT B	B uGcuGGAcAuGAGAuGGAGTT B	B GAGGcuAcAGGGGuuAGccTT B	B GAccuAccucAAAGGGCAGTT B	cAuuAAGcAccAcAGcAuATsT	cuccAucucAuGuccAGcATsT	GG 1A Account A Good Class	H	cuGcccuuuGAGGuAGGuc181	UAUGCUGUGGUGCUUAAUGTT	UGCUGGACAUGAGAUGGAGTT	GAGGCUACAGGGGUUAGCCTT	GACCUACCUCAAAGGGCAGTT	CAUUAAGCACCACAGCAUALI	CUCCAUCUCAUGUCCAGCATT	TTOILOGORISTICATION	GGCUAACCCCUGUAGCCUCT
(146C)	16 RelA:308L21 siRNA		045C) 08 RelA:1975L21 siRNA	31308 RelA:1957U21 siRNA	31309 RelA:1975L21 siRNA		21 RELA:1975L21 siRNA (1957C) inv stah05		30874 SCD:2520U21 siRNA	30875 SCD:3785U21 siRNA	30876 SCD:4774U21 siRNA			(2520C) stab05	30879   SCD:3803LZ1 SIRIVA (3785C) stab05	30880 SCD:4792L21 siRNA (4774C) stab05			31023 SCD:3785U21 siRNA	31024 SCD:4774U21 siRNA	31097 SCD:1013L21 siRNA	(995C) 31098 SCD:2538L21 siRNA	(2520C)	31099 SCD:3803L21 siRNA
	antisense 31106	antisense 31107	antisense 31108	sense 3130	antisense 3130	sense 31320	antisense 31321	sense 30873	sense 308	sense 308	sense 308	antisense 30877	antisense 30878		antisense   308	antisense 308	sense 31021	$\top$	1-	T	Ø	anticence 310		antisense 310
	162 a	163 a	164 a	164	164 a	164	164 8	165	166	167	168	165	166	7	167	168	165	166	167	168	165	186	3	167
	GAUGGCUUCUAUGAGGCUGAGCU	UGUGUGACAAGGUGCAGAAAGAG	UCCUCCAGCUUCUGGUACUCUCC	UCCUCCAGCUUCUGGUACUCUCC	Uccuccaecuucueeuacucucc	UCCUCCAGCUUCUGGUACUCUCC	uccuccaecuucueeuacucucc	GAUAUGCUGUGGUGCUUAAUGCC	ACUGCUGGACAUGAGAUGGAGAG	UAGAGGCUACAGGGGUUAGCCUG	CUGACCUACCUCAAAGGGCAGUU	GAUAUGCUGUGGUGCUUAAUGCC			UAGAGGCUACAGGGGUUAGCCUG	CUGACCUACCUCAAAGGGCAGUU	229118411123113311311331141143	GAUAUGCUGUGGGGGC			GAUAUGCUGUGGUGCU	VO VOI IVO VOI IVO	ACUGCUGGACAUGAGAGGAGAG	UAGAGGCUACAGGGGUUAGCCUG
-	288	643	1955	1955	1955	1955	1955	993	2518	3783	4772	993	2518	2	3783	4772					993		2518	3783
	RELA	RELA	RELA	RELA	RELA	RELA	RELA	SCD	SCD	SCD	SCD	SCD	200	3	SCD	SCD			SCD	G) (8		3	SCD	SCD

874

875

853 854 856 856 857 858 858

860

861

1244

TERT

641

TERT

2495

TERT

1136

TERT

1790 AGUGUCU

TERT

2915

TERT

2994 UGAAGUGI

TERT

1136 GUGGAGA

TERT

1790 AGUGUCU

TERT

2915

TERT

2994

TERT

862

863

864

865

998

867

868

869

870

872

871

1526 AGGGAUAA

TGFB1

2383

TGFB1

2484

TGFB1

2566

TGFB1

1526 AGGGAUAA

TGFB1

873

	CUGCCCUUUGAGGUAGGUCTT	GCGCACGUGGGAAGCCCUGGC	CAGAGGCUGUGCGAGCGCGGC	UCUGGGAUGCGAACGGGCCUG	UGGGAACCACGCGCAGUGCCC	CCACCACGCGUGCGCAUCAG	CAGGGCUUCCCACGUGCGCAG	CGCGCUCGCACAGCCUCUGCA	GGCCCGUUCGCAUCCCAGACG	GCACUGCGCGUGGUUCCCAAG	GAUGCGCACGGCGUGGUGGCA	B GGAGAccAucuuucuGGGuTT B	B uGucuGGAGcAAGuuGcAATT B	B cAGAGccAGucucAccuucTT B	B AAGuGucAcAGccuGuuucTT B	AcccAGAAAGAuGGucuccTsT	uuGcAAcuuGcuccAGAcATsT	GAAGGuGAGAcuGGcucuGTsT	GAAAcAGGcuGuGAcAcuuTsT	B GGAUAAcAcAcuGcAAGuGTT B	B AuAGcAAcacucuGAGAuGTT B	B AccuGcuuuAGuGGGGGAuTT B	B GcAcuuuuGGGAGGCAGAGTT B	cAcuuGcAGuGuGuuAuccTsT
	antisense   31100   SCD:4792L21 siRNA   (4774C)	29960 TERT:19U21 siRNA	29961 TERT:311U21 siRNA	29962 TERT:643U21 siRNA	29963 TERT:1246U21 siRNA	29964 TERT:2497U21 siRNA	29965 TERT:39L21 siRNA (19C)	antisense 29966 TERT:331L21 siRNA (311C)	antisense 29967 TERT:663L21 siRNA (643C)			4 <b>–</b> 0.		TERT:2917U21 siRNA stab04	TERT:2996U21 siRNA stab04	TERT:1156L21 siRNA (1138C) stab05	TERT:1810L21 siRNA (1792C) stab05	TERT:2935L21 siRNA (2917C) stab05	TERT:3014L21 siRNA (2996C) stab05	TGFb:1528U21 siRNA stab04	TGFb:2385U21 siRNA stab04	30883 TGFb:2486U21 siRNA stab04	30884 TGFb:2568U21 siRNA stab04	antisense 30885 TGFb:1546L21 siRNA
00770	31100	29960	29961	29962	29963	29964		29966	29967	29968	29969	30905	30906	30907	30908	60608	30910	30911	30912	30881	30882	30883	30884	30885
	antisense	seuse	sense	sense	sense	sense	antisense	antisense	antisense	antisense	antisense	sense	sense	sense	seuse	antisense	antisense	antisense	antisense	sense	seuse	seuse	seuse	antisense
90,	168	169	170	171	172	173	169	170	171	172	173	174	175	176	177	174	175	176	177	178	179	180	181	178
-	Z COGACCOACCOCAGGGGCAGUO	CUGCGCACGUGGGAAGCCCUGGC	UGCAGAGGCUGUGCGAGCGCGGC	CGUCUGGGAUGCGAACGGGCCUG	4 CUUGGGAACCACGCGCAGUGCCC	5 UGCCACCACGCGUGCGCAUCAG	CUGCGCACGUGGGAAGCCCUGGC	UGCAGAGGCUGUGCGAGCGCGGC	CGUCUGGGAUGCGAACGGGCCUG	CUUGGGAACCACGCGCAGUGCCC	) UGCCACCACGCGUGCGCAUCAG	GUGGAGACCAUCUUCUGGGUUC	AGUGUCUGGAGCAAGUUGCAAAG	AUCAGAGCCAGUCUCACCUUCAA	UGAAGUGUCACAGCCUGUUUCUG	GUGGAGACCAUCUUCUGGGUUC	AGUGUCUGGAGCAAGUUGCAAAG	AUCAGAGCCAGUCUCACCUUCAA	UGAAGUGUCACAGCCUGUUUCUG	AGGGAUAACACACUGCAAGUGGA	CCAUAGCAACACUCUGAGAUGGC	GAACCUGCUUNAGUGGGGGAUAG	UAGCACUUUUGGGAGGCAGAGAU	AGGGAUAACACACUGCAAGUGGA

4772

SCD

309 641 1244 2495

TERT TERT TERT TERT TERT

	876	877	878	T 879	T 880								3 888	B 883	В 890					71 895 71	1	-	TT:
	cAucucAGAGuGuuGcuAuTsT	AuccccAcuAAAGcAGGuTsT	cucuGccucccAAAAGuGcTsT	GGAUAACACACUGCAAGUGTT	AUAGCAACACUCUGAGAUGTT	ACCUGCUUNAGUGGGGGAUT	GCACUUUUGGGAGGCAGAGTT	CACUUGCAGUGUGUUAUCCTT	CAUCUCAGAGUGUUGCUAUTI	AUCCCCACUAAAGCAGGUTT	CUCUGCCUCCCAAAAGUGCTT	B GGAcAccAuGAGcAcuGAATT B	B GuuccucAGccucuucuccTT B	B ccuAccAGAccAAGGucAATT B	B AGGccuuccucuccAGATT B	uucAGuGcucAuGGuGuccTsT	GGAGAAGAGGcuGAGGAAcTsT	uuGAccuuGGucuGGuAGGTsT	ucuGGAGAGAGGAAGGccuTsT	GGACACCAUGAGCACUGAATT	TTAACITOOAACCACACACACACACACACACACACACACACACACA	AGGCIIIOCIICOCAGAGGATT	TION DE LO COLLOYOR D
(4E28C) stab05				2206C) Staboo	31033 1GFD: 1320021 SILVIN	31055 TGFb:2486U21 siRNA	31056 TGFb:2568U21 siRNA	antisense 31129 TGFb:1546L21 siRNA	1	<u> </u>		30889 TNFa:79U21 siRNA	90 TNFa:178U21 siRNA	91 TNFa:570U21 siRNA	92 TNFa:1152U21 siRNA	93 TNFa:97L21 siRNA (79C)	94 TNFa:196L21 siRNA (178C) stab05	30895 TNFa:588L21 siRNA (570C) stab05		31408 TNFa:79U21 siRNA	31409 TNFa:178UZ1 SIKINA	31410 TNFa:570U21 siRNA	sense 31411
	antisense 30886	antisense 30887	antisense 30888	$\top$		sense 3105	T	antisense 3112	antisense 31130	antisense 31131	antisense 31132	sense 308	sense 30890	sense 30891	sense 30892	antisense 30893	antisense 30894	antisense 308	antisense 30896		sense 314		sense 314
	179	180	181	7,1	1/8	2 6	2 2	<del>                                     </del>	179	180	181	182	183	184	185	182	183	184	185	182	183	184	185
	CCAUAGCAACACUCUGAGAUGGC	GAACCUGCUUUAGUGGGGGAUAG	UAGCACUUUUGGGAGGCAGAGAU			CCAUAGCAACACUCUGAGAUGGC			CCAUAGCAACACUCUGAGAUGGC	GAACCUGCUUNAGUGGGGGAUAG	UAGCACUUUUGGGAGGCAGAGAU	AAGGACACCAUGAGCACUGAAAG	UNGUNCCUCAGCCUCUNCUCCUN	CUCCUACCAGACCAAGGUCAACC	UNAGGCCUUCCUCUCCAGAUG	AAGGACACCAUGAGCACUGAAAG	UUGUUCCUCAGCCUCUUCCUU	CUCCUACCAGACCAAGGUCAACC	UUAGGCCUUCCUCUCCAGAUG	AAGGACACCAUGAGCACUGAAAG	-	+	UNAGGCCUUCCUCUCCAGAUG
	2383	2484	2566			_	2484		2383	2484	2566	77	176	568	1150	77	176	268	1150	11	176	268	1150
	TGFB1	TGFB1	TGFB1		TGFB1	TGFB1	TGFB1	TGFB1	TGFB1	TGFB1	TGFB1	TNF	TNF	ŢNT	TNT FNT	F.	TNF	AN THE	ANT.	F F	AN T	上	HAT.

900	901	805	
GGAGAAGAGGCUGAGGAACTT	UUGACCUUGGUCUGGUAGGTT	UCUGGAGAGAGGCCUTT	
JCUCCUU 183 antisense 31413 TNFa:196L21 siRNA (178C)	TNFa:588L21 siRNA (570C)	TNFa:1170L21 siRNA	(1152C)
31413	31414	31415	
antisense	antisense	antisense	
183	184	185	
TNF 176 UUGUUCCUCAGCCUCUUCUCCUU	TNF 568 CUCCUACCAGACCAAGGUCAACC 184 antisense 31414 TNFa:588L21 siRNA (570C)	TNF 1150 UUAGGCCUUCCUCCUCCAGAUG 185 antisense 31415 TNFa:1170L21 siRNA	
176	268	1150	
TNF	TNF	TNF	

Uppercase = ribonucleotide u,c = 2'-deoxy-2'-fluoro U,C T = thymidine B = inverted deoxy abasic s = phosphorothioate linkage A = deoxy Adenosine G = deoxy Guanosine

## Table II

A. 2.5 μmol Synthesis Cycle ABI 394 Instrument

Reagent	Equivalents	Amount	Wait Time* DNA	Wait Time* 2'-O-methyl	Wait Time*RNA
Phosphoramidites	6.5	163 µL	45 sec	2.5 min	7.5 min
S-Ethyl Tetrazole	23.8	238 µL	45 sec '	2.5 min	7.5 min
Acetic Anhydride	100	233 µL	5 sec	5 sec	5 sec
N-Methyl Imidazole	186	233 µL	5 sec	5 sec	5 sec
TCA	176	2.3 mL	21 sec	21 sec	21 sec
Iodine	11.2	1.7 mL	.45 sec	45 sec	45 sec
Beaucage	12.9	645 µL	100 sec	300 sec	300 sec
Acetonitrile	NA	6.67 mL	NA	NA	NA

B. 0.2 µmol Synthesis Cycle ABI 394 Instrument

Reagent	Equivalents	Amount	Wait Time* DNA	Wait Time* 2'-O-methyl	Wait Time*RNA
Phosphoramidites	15	31 µL	45 sec	233 sec	465 sec
S-Ethyl Tetrazole	38.7	31 µL	45 sec	233 min	465 sec
Acetic Anhydride	655	124 µL	5 sec	5 sec	5 sec
N-Methyl Imidazole	1245	124 µL	5 sec	5 sec	5 sec
TCA	700	732 µL	10 sec	10 sec	10 sec
Iodine	20.6	244 μL	15 sec	15 sec	15 sec
Beaucage	7.7	232 µL	100 sec	300 sec	300 sec
Acetonitrile	NA	2.64 mL	NA	NA	NA

C.  $0.2\,\mu mol$  Synthesis Cycle 96 well Instrument

Reagent	Equivalents:DNA/ 2'-O-methyl/Ribo	Amount: DNA/2'-O- methyl/Ribo	Wait Time* DNA	Wait Time* 2'-O- methyl	Wait Time* Ribo
Phosphoramidites	22/33/66	40/60/120 μL	60 sec	180 sec	360sec
S-Ethyl Tetrazole	70/105/210	40/60/120 μL	60 sec	180 min	360 sec
Acetic Anhydride	265/265/265	50/50/50 μL	10 sec	10 sec	10 sec
N-Methyl Imidazole	502/502/502	50/50/50 μL	10 sec	10 sec	10 sec
TCA	238/475/475	250/500/500 μL	15 sec	15 sec	15 sec
lodine	6.8/6.8/6.8	80/80/80 µL	30 sec	30 sec	30 sec
Beaucage	34/51/51	80/120/120	100 sec	200 sec	200 sec
Acetonitrile	NA	1150/1150/1150 µL	NA	NA	NA

- Wait time does not include contact time during delivery.
- Tandem synthesis utilizes double coupling of linker molecule

Table III

Filter (1.0	Group	Solution on	Stock VEGF	Number	Injectate	Dose	Conc.
L     NA   S   water   NA   NA   NA   NA   NA   NA   NA   N		Filter (1.0	concentration	of			
1 Tris-Cl pH 6.9 2 R&D Systems VEGF-carrier free 75 μM 3 R&D Systems VEGF-carrier free 75 μM 4 R&D Systems VEGF-carrier free 75 μM 5 R&D Systems VEGF-carrier free 75 μM 5 R&D Systems VEGF-carrier free 75 μM 5 R&D Systems VEGF-carrier free 75 μM 6 R&D Systems VEGF-carrier free 75 μM 7 R&D Systems VEGF-carrier free 75 μM 6 R&D Systems VEGF-carrier free 75 μM 7 R&D Systems VEGF-carrier free 75 μM 8 R&D Systems VEGF-carrier free 75 μM 7 R&D Systems VEGF-carrier free 75 μM 8 R&D Systems VEGF-carrier free 75 μM 8 R&D Systems VEGF-carrier free 75 μM  8 R&D Systems VEGF-carrier free 75 μM  8 R&D Systems VEGF-carrier free 75 μM  8 R&D Systems VEGF-carrier free 75 μM  8 R&D Systems VEGF-carrier free 75 μM  8 R&D Systems VEGF-carrier free 75 μM  8 R&D Systems VEGF-carrier free 75 μM  8 R&D Systems VEGF-carrier free 75 μM  8 R&D Systems VEGF-carrier free 75 μM  8 R&D Systems VEGF-carrier free 75 μM  8 R&D Systems VEGF-carrier free 75 μM  8 R&D Systems VEGF-carrier free 75 μM  8 R&D Systems VEGF-carrier free 75 μM  8 R&D Systems VEGF-carrier free 75 μM  8 R&D Systems VEGF-carrier free 75 μM  8 R&D Systems VEGF-carrier free 75 μM  9 1 0.167 μg/eye μg/μL  1 0.167 μg/eye γμg/μL		μL)		Animals			
Color					3,713		
2 R&D Systems VEGF-carrier free 75 μM  3 R&D Systems VEGF-carrier free 75 μM  4 R&D Systems VEGF-carrier free 75 μM  5 Site 2340 10 μg/eye μg/μL siRNA  6 R&D Systems VEGF-carrier free 75 μM  6 R&D Systems VEGF-carrier free 75 μM  7 R&D Systems VEGF-carrier free 75 μM  7 R&D Systems VEGF-carrier free 75 μM  8 R&D Systems VEGF-carrier free Stab1 siRNA	1	Tris-Cl pH	NA	5	water	NA	NA
VEGF-carrier free   75 μM   3.53 μg/μL   5   Site 2340   10   1.67							
Site 2340   10   1.67   μg/eye   μg/μL   5   Site 2340   μg/eye   Site 2340   μg/eye   μg/μL   5	2		3.53 μg/μL	5	water	NA	NA
75 μM   3   R&D Systems   3.53 μg/μL   5   Site 2340   10   1.67   μg/eye   μg/μL   5   Site 2340   μg/eye   μg/μL   5   Site 2340   μg/eye   μg/μL   5   Site 2340   3   μg/eye   μg/μL   5   Site 2340   3   μg/eye   μg/μL   5   Site 2340   μg/eye   μg/μL   5		VEGF-carrier					
3   R&D Systems   VEGF-carrier   free   75 μM     3.53 μg/μL   5   Site 2340   μg/eye   μg/μL     1.67 μg/eye   1.67 μg/eye   μg/μL     1.67 μg/eye   1.67 μg		free					
VEGF-carrier free   Stab1 siRNA   μg/eye   μg/μL							
free   75 μM   3.53 μg/μL   5   Site 2340   3   μg/eye   μg/μL   5   Site 2340   3   μg/eye   μg/μL   5   Site 2340   3   μg/eye   μg/μL   5   Site 2340   1   μg/eye   μg/μL   5   Site 2340   Site 2340   Site 2340   Site 2340   μg/eye   μg/μL   5   Site 2340   Sit	3		3.53 μg/μL	5	Site 2340	10	1.67
75 μM   3.53 μg/μL   5   Site 2340   3   μg/eye   μg/μL     5   R&D Systems   3.53 μg/μL   5   Site 2340   μg/eye   μg/μL     5   R&D Systems   3.53 μg/μL   5   Site 2340   1   μg/eye   μg/μL     6   R&D Systems   75 μM   5   Inactive   Site 2340   μg/eye   μg/μL     6   R&D Systems   3.53 μg/μL   5   Inactive   Site 2340   μg/eye   μg/μL     7   R&D Systems   3.53 μg/μL   5   Inactive   Site 2340   μg/eye   μg/μL     7   R&D Systems   3.53 μg/μL   5   Inactive   Site 2340   μg/eye   μg/μL     7   R&D Systems   3.53 μg/μL   5   Inactive   Site 2340   μg/eye   μg/μL     8   R&D Systems   Stab1   SiRNA     8   R&D Systems   Stab1   SiRNA   Site 2340   μg/eye   μg/μL     8   R&D Systems   Stab1   Site 2340   μg/eye   μg/μL     8   R&D Systems   Stab1   Site 2340   μg/eye   μg/μL     9   Stab1   Site 2340   μg/eye   μg/μL     9   Site 2340   μg/eye   μg/μL     9   Site 2340   μg/eye   μg/μL     1   O.167   μg/eye   μg						μg/eye	μg/μL
4 R&D Systems VEGF-carrier free 75 μΜ  5 R&D Systems VEGF-carrier free 75 μΜ  6 R&D Systems VEGF-carrier free 75 μΜ  7 R&D Systems VEGF-carrier free 75 μΜ  7 R&D Systems VEGF-carrier free 75 μΜ  8 R&D Systems VEGF-carrier free 75 μβ/μL  8 R&D Systems VEGF-carrier free 75 μβ/μL  8 R&D Systems VEGF-carrier free 75 μβ/μL		_			siRNA		_
VEGF-carrier free   Stab1 siRNA   μg/eye μg/μL     5   R&D Systems   3.53 μg/μL   5   Site 2340   1 μg/eye μg/μL     6   R&D Systems   3.53 μg/μL   5   Inactive   10 μg/eye   μg/μL     6   R&D Systems   3.53 μg/μL   5   Inactive   Stab1   siRNA     75 μM   7   R&D Systems   3.53 μg/μL   5   Inactive   Stab1   siRNA     7   R&D Systems   3.53 μg/μL   5   Inactive   3 μg/eye   μg/μL     75 μM   Site 2340   Site 2340   μg/eye   μg/μL     75 μM   Site 2340   Site 2340   μg/eye   μg/μL     8   R&D Systems   3.53 μg/μL   5   Inactive   Site 2340   μg/eye   μg/μL     8   R&D Systems   VEGF-carrier   free   Site 2340   Site 2340   Site 2340   μg/eye   μg/μL     8   R&D Systems   Site 2340   μg/eye   μg/μL     8   R&D Systems   Site 2340   Site 2		<del></del>					
free   75 μM   5   Site 2340   1   μg/eye   μg/μL   5   Site 2340   Site 2340   μg/eye   μg/μ	4	•	3.53 μg/μL	5	1	3	0.5
75 μM   3.53 μg/μL   5   Site 2340   1   μg/eye   μg/μL   5   Site 2340   1   μg/eye   μg/μL   5   Site 2340   1   μg/eye   μg/μL   5   Site 2340   μg/eye   Site 2340   μg/eye   Site 2340   μg/eye   μg/μL   5   Site 2340   μg/eye   Site 2340   μg/eye   μg/μL   5   Site 2340   μg/eye   Site 2340   μg/eye   μg/μL   5   Site 2340   μg/eye   μg/μL   Site 2340   μg/e					i	μg/eye	μg/μL
Site 2340   1					siRNA		
VEGF-carrier free							
free   siRNA   siRNA   free   siRNA   free   siRNA   free   siRNA   free   sith   free   free   sith   free   free   free   free   sith   free   f	5	<b>-</b>	3.53 μg/μL	5	i	_	
75 μM		i			i	μg/eye	μg/μL
6 R&D Systems VEGF-carrier free $75 \mu M$ 7 R&D Systems VEGF-carrier free $75 \mu M$ 8 R&D Systems VEGF-carrier Site 2340 $97 \mu M$ 8 R&D Systems VEGF-carrier Site 2340 $97 \mu M$ 8 Stab1  8 Stab1  9 Stab1  9 O.167  9 Site 2340 $97 \mu M$					siRNA		
VEGF-carrier free free			2.52 / 1	<u> </u>			
free   Stab1   siRNA	0		3.33 μg/ μL	5	1		
75 μM   siRNA     siRNA						μg/eye	μg/μL
7R&D Systems VEGF-carrier free 75 μΜ3.53 μg/μL5Inactive Site 2340 Stab1 siRNA30.5 μg/μL8R&D Systems VEGF-carrier free3.53 μg/μL5Inactive Stab110.167 μg/eyeVEGF-carrier freeStab15Inactive Site 2340 Stab1							
VEGF-carrier   free   free   75 μM   Site 2340   μg/eye   μg/μL     μg/eye   μg/μL     μg/eye   μg/μL     μg/eye   μg/μL     μg/eye   μg/μL     1   0.167     μg/eye   μg/μL     1   μg/eye   μg/μL     μg/eye	7		3 53 ug/ut	<u> </u>		2	0.5
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$			υ.υυ μg/ μL	3			
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$		•				μg/eye	μg/μL
8 R&D Systems $VEGF$ -carrier free $Site 2340$ μg/eye $VegF$ -carrier $VegF$ -carr		`					
VEGF-carrier free Stab1	8	<del></del>	3.53 ug/uI	5		1	0.167
free Stab1			0.00 MB/ ME				
						μg/ cye	нв/ нь
I CHIVE I SIKINA I I		75 μM			siRNA		

**Table IV** 

Non-limiting examples of Stabilization Chemistries for chemically modified siNA constructs

Chemistry	pyrimidine	Purine	сар	p=S	Strand
"Stab 1"	Ribo	Ribo	-	5 at 5'-end 1 at 3'-end	S/AS
"Stab 2"	Ribo	Ribo	-	All linkages	Usually AS
"Stab 3"	2'-fluoro	Ribo	-	4 at 5'-end 4 at 3'-end	Usually S
"Stab 4"	2'-fluoro	Ribo	5' and 3'- ends	1	Usually S
"Stab 5"	2'-fluoro	Ribo	-	1 at 3'-end	Usually AS
"Stab 6"	2'-O-Methyl	Ribo	5' and 3'- ends	_	Usually S
"Stab 7"	2'-fluoro	2'-deoxy	5' and 3'- ends	-	Usually S
"Stab 8"	2'-fluoro	2'-O- Methyl	-	1 at 3'-end	Usually AS
"Stab 9"	Ribo	Ribo	5' and 3'- ends	-	Usually S
"Stab 10"	Ribo	Ribo	-	1 at 3'-end	Usually AS
"Stab 11"	2'-fluoro	2'-deoxy		1 at 3'-end	Usually AS

<sup>5</sup> CAP = any terminal cap, see for example Figure 10.

All Stab 1-11 chemistries can comprise 3'-terminal thymidine (TT) residues

All Stab 1-11 chemistries typically comprise 21 nucleotides, but can vary as described herein.

S = sense strand

10 AS = antisense strand

Table V

Acc#	Description
NM 002825	Homo sapiens pleiotrophin (heparin binding growth factor 8, neurite growth-
- 12.1_002020	promoting factor 1) (PTN), mRNA
NM 033418	Homo sapiens hypothetical protein MGC9084 (MGC9084), mRNA
NM 033111	Homo sapiens LOC88523 (LOC88523), mRNA
NM 032564	Homo sapiens diacylglycerol O-acyltransferase homolog 2 (mouse) (DGAT2),
	mRNA
NM_032311	Homo sapiens KIAA1649 protein (KIAA1649), mRNA
NM_022130	Homo sapiens golgi phosphoprotein 3 (coat-protein) (GOLPH3), mRNA
NM_021980	Homo sapiens optineurin (OPTN), mRNA
NM_000660	Homo sapiens transforming growth factor, beta 1 (Camurati-Engelmann disease) (TGFB1), mRNA
NM_020423	Homo sapiens hypothetical protein LOC57147 (LOC57147), mRNA
NM_020351	Homo sapiens smooth muscle cell-expressed and macrophage conditioned
	medium-induced protein smag-64 (LOC57086), mRNA
NM_019556	Homo sapiens hypothetical protein dJ473B4 (DJ473B4), mRNA
NM_018676	Homo sapiens TMTSP for transmembrane molecule with thrombospondin
	module (LOC55901), mRNA
NM_016265	Homo sapiens GIOT-3 for gonadotropin inducible transcription repressor-3
	(GIOT-3), mRNA
NM_016531	Homo sapiens Kruppel-like factor 3 (basic) (KLF3), mRNA
NM_016372	Homo sapiens seven transmembrane domain orphan receptor (TPRA40), mRNA
NM_016211	Homo sapiens yeast Sec31p homolog (KIAA0905), mRNA
NM_014933	Homo sapiens yeast Sec31p homolog (KIAA0905), mRNA
NM_014706	Homo sapiens squamous cell carcinoma antigen recognised by T cells 3 (SART3), mRNA
NM 014463	Homo sapiens Lsm3 protein (LSM3), mRNA
NM_014288	Homo sapiens integrin beta 3 binding protein (beta3-endonexin) (ITGB3BP),
_	mRNA
NM_013443	Homo sapiens CMP-NeuAC:(beta)-N-acetylgalactosaminide (alpha)2,6-
	sialyltransferase member VI (VI), mRNA
NM_012404	Homo sapiens pp32 related 2 (PP32R2), mRNA
NM_012403	Homo sapiens pp32 related 1 (PP32R1), mRNA
NM_006710	Homo sapiens COP9 homolog (COP9), mRNA
NM_006117	Homo sapiens peroxisomal D3,D2-enoyl-CoA isomerase (PECI), mRNA
NM_005839	Homo sapiens serine/arginine repetitive matrix 1 (SRRM1), mRNA
NM_004264	Homo sapiens SRB7 suppressor of RNA polymerase B homolog (yeast)
	(SURB7), mRNA
NM_003714	Homo sapiens stanniocalcin 2 (STC2), mRNA
NM_003122	Homo sapiens serine protease inhibitor, Kazal type 1 (SPINK1), mRNA
NM_003690	Homo sapiens protein kinase, interferon-inducible double stranded RNA
	dependent activator (PRKRA), mRNA
NM_015526	Homo sapiens CLIP-170-related protein (CLIPR-59), mRNA
NM_033401	Homo sapiens cell recognition protein CASPR4 (CASPR4), mRNA
NM_023037	Homo sapiens hypothetical protein CG003 (13CDNA73), mRNA
NM_021817	Homo sapiens brain link protein-1 (BRAL1), mRNA
NM_016222	Homo sapiens DEAD-box protein abstrakt (ABS), mRNA
NM_003744	Homo sapiens numb homolog (Drosophila) (NUMB), mRNA
NM_032682	Homo sapiens forkhead box P1 (FOXP1), mRNA
NM_003681	Homo sapiens pyridoxal (pyridoxine, vitamin B6) kinase (PDXK), mRNA

	1 1:100
NM_001685	Homo sapiens ATP synthase, H+ transporting, mitochondrial F0 complex, subunit F6 (ATP5J), mRNA
NM_017954	Homo sapiens hypothetical protein FLJ20761 (FLJ20761), mRNA
NM 015626	Homo sapiens SOCS box-containing WD protein SWiP-1 (WSB1), mRNA
NM 130795	Homo sapiens regulator of G-protein signalling 3 (RGS3), mRNA
NM 030877	Homo sapiens chromosome 20 open reading frame 33 (C20orf33), mRNA
NM 080830	Homo sapiens cystatin 11 (CST11), mRNA
NM 032329	Homo sapiens p28 ING5 (ING5), mRNA
NM 022917	Homo sapiens nucleolar RNA-associated protein (Nrap), mRNA
	Homo sapiens adaptor-related protein complex 2, alpha 1 subunit (AP2A1),
NM_130787	mRNA
NM_024744	Homo sapiens (ALS2CR8), mRNA
NM_018984_	Homo sapiens slingshot 1 (hSSH-1), mRNA
NM_106552	Homo sapiens hypothetical protein FLJ14249 similar to HS1 binding protein 3
	(FLJ14249), transcript variant 2, mRNA
NM 022460	Homo sapiens hypothetical protein FLJ14249 similar to HS1 binding protein 3
	(FLJ14249), transcript variant 1, mRNA
NM 130446	Homo sapiens kelch-like protein KLHL6 (KLHL6), mRNA
NM 020314	Homo sapiens esophageal cancer associated protein (MGC16824), mRNA
NM_130395	Homo sapiens Werner helicase interacting protein (WHIP), transcript variant 2, mRNA
NM_020135	Homo sapiens Werner helicase interacting protein (WHIP), transcript variant 1, mRNA
NM 130388	Homo sapiens ankyrin repeat and SOCS box-containing 12 (ASB12), mRNA
NM 130387	Homo sapiens ankyrin repeat and SOCS box-containing 14 (ASB14), mRNA
NM 007191	Homo sapiens WNT inhibitory factor 1 (WIF1), mRNA
NM 052950	Homo sapiens WD40- and FYVE-domain containing protein 2 (WDF2), mRNA
NM_025042	Homo sapiens Williams-Beuren syndrome chromosome region 23 (WBSCR23), mRNA
NM_080706	Homo sapiens transient receptor potential cation channel, subfamily V, member 1 (TRPV1), transcript variant 3, mRNA
NM_080705	Homo sapiens transient receptor potential cation channel, subfamily V, member 1 (TRPV1), transcript variant 4, mRNA
NM_080704	Homo sapiens transient receptor potential cation channel, subfamily V, member 1 (TRPV1), transcript variant 1, mRNA
NM_018727	Homo sapiens transient receptor potential cation channel, subfamily V, member 1 (TRPV1), transcript variant 2, mRNA
NM_080879	Homo sapiens SOCS box containing protein RAR2A (RAR2A), mRNA
NM 080871	Homo sapiens ankyrin repeat and SOCS box-containing 10 (ASB10), mRNA
NM 080870	Homo sapiens DPCR1 protein (DPCR1), mRNA
NM 080870	Homo sapiens chromosome 20 open reading frame 152 (C20orf152), mRNA
NM 080829	Homo sapiens chromosome 20 open reading frame 175 (C20orf175), mRNA
NM 080828	Homo sapiens chromosome 20 open reading frame 173 (C20orf173), mRNA
	Homo sapiens G protein-coupled receptor 78 (GPR78), mRNA
NM 080819	Homo sapiens of protein-coupled receptor 78 (OFR/8), interval  Homo sapiens chromosome 20 open reading frame 164 (C20orf164), mRNA
NM_080752	Homo sapiens chromosome 20 open reading frame 163 (C20orf163), mRNA
NM_080749	
NM_080745	Homo sapiens ring finger protein 36 (RNF36), mRNA
NM_080738	Homo sapiens EDAR-associated death domain (EDARADD), mRNA
NM_014970	Homo sapiens kinesin-associated protein 3 (KIFAP3), mRNA
NM_021058	Homo sapiens H2B histone family, member R (H2BFR), mRNA
NM_021064	Homo sapiens H2A histone family, member P (H2AFP), mRNA
NM_080491	Homo sapiens GRB2-associated binding protein 2 (GAB2), transcript variant 1,

	mRNA
ND4 012206	Homo sapiens GRB2-associated binding protein 2 (GAB2), transcript variant 2,
NM_012296	-
) D ( 007047	mRNA Homo sapiens AP1 gamma subunit binding protein 1 (AP1GBP1), transcript
NM_007247	
-	variant 1, mRNA
NM_080551	Homo sapiens AP1 gamma subunit binding protein 1 (AP1GBP1), transcript
	variant 3, mRNA
NM_080550	Homo sapiens AP1 gamma subunit binding protein 1 (AP1GBP1), transcript
	variant 2, mRNA
NM_000982	Homo sapiens ribosomal protein L21 (RPL21), mRNA
NM_003913	Homo sapiens serine/threonine-protein kinase PRP4 homolog (PRP4), mRNA
NM_002475	Homo sapiens myosin light chain 1 slow a (MLC1SA), mRNA
NM_002729	Homo sapiens hematopoietically expressed homeobox (HHEX), mRNA
NM 005893	Homo sapiens calicin (CCIN), mRNA
NM 017593	Homo sapiens homolog of mouse BMP-2 inducible kinase (BIKE), mRNA
NM 032027	Homo sapiens beta-amyloid binding protein precursor (BBP), mRNA
NM 004051	Homo sapiens 3-hydroxybutyrate dehydrogenase (heart, mitochondrial) (BDH),
1111_001001	nuclear gene encoding mitochondrial protein, mRNA
NM 006576	Homo sapiens advillin (AVIL), mRNA
NM 013375	Homo sapiens TATA-binding protein-binding protein (ABT1), mRNA
NM 058219	Homo sapiens homolog of yeast mRNA transport regulator 3 (MTR3), mRNA
NM 058237	Homo sapiens HEAT-like repeat-containing protein (KIAA1622), transcript
NWI_030237	variant 1, mRNA
NM 020958	Homo sapiens HEAT-like repeat-containing protein (KIAA1622), transcript
NWI_020936	variant 2, mRNA
NM 004702	Homo sapiens cyclin E2 (CCNE2), transcript variant 3, mRNA
	Homo sapiens cyclin E2 (CCNE2), transcript variant 1, mRNA
NM_057749	Homo sapiens cyclin E2 (CCNE2), transcript variant 1, incova  Homo sapiens cyclin E2 (CCNE2), transcript variant 2, mRNA
NM_057735	
NM_002013	Homo sapiens FK506 binding protein 3 (25kD) (FKBP3), mRNA
NM_004724	Homo sapiens ZW10 homolog, centromere/kinetochore protein (Drosophila)
37.6.0551.50	(ZW10), mRNA
NM_057159	Homo sapiens endothelial differentiation, lysophosphatidic acid G-protein-
	coupled receptor, 2 (EDG2), transcript variant 2, mRNA
NM_001401	Homo sapiens endothelial differentiation, lysophosphatidic acid G-protein-
	coupled receptor, 2 (EDG2), transcript variant 1, mRNA
NM_015084	Homo sapiens mitochondrial ribosomal protein S27 (MRPS27), nuclear gene
	encoding mitochondrial protein, mRNA
NM_033281	Homo sapiens mitochondrial ribosomal protein S36 (MRPS36), nuclear gene
	encoding mitochondrial protein, mRNA
NM_005830	Homo sapiens mitochondrial ribosomal protein S31 (MRPS31), nuclear gene
	encoding mitochondrial protein, mRNA
NM_012062	Homo sapiens dynamin 1-like (DNM1L), transcript variant 1, mRNA
NM_005648	Homo sapiens transcription elongation factor B (SIII), polypeptide 1 (15kD,
	elongin C) (TCEB1), mRNA
NM_007070	Homo sapiens FKBP-associated protein (FAP48), transcript variant 2, mRNA
NM 053274	Homo sapiens FKBP-associated protein (FAP48), transcript variant 1, mRNA
NM_054113	Homo sapiens DNA-dependent protein kinase catalytic subunit-interacting
	protein 3 (KIP3), mRNA
NM 003726	Homo sapiens src family associated phosphoprotein 1 (SCAP1), mRNA
NM 012308	Homo sapiens F-box and leucine-rich repeat protein 11 (FBXL11), mRNA
NM_030913	Homo sapiens sema domain, transmembrane domain (TM), and cytoplasmic
14147_020212	domain, (semaphorin) 6C (SEMA6C), mRNA
	domain, (Semaphorin) oc (SEM 100), interio

PCT/US03/05028 WO 03/074654

M 021163	Homo sapiens RB-associated KRAB repressor (RBAK), mRNA
M 033632	Homo sapiens F-box and WD-40 domain protein / (archipelago homolog,
(1,1	Descentile) (FRYW7) transcript variant 1, mKNA
IM_018315	Homo saniens F-box and WD-40 domain protein / (archipelago nonlolog,
111_010515	Drosophila) (FBXW7), transcript variant 2, mRNA
JM 012168	Home conjens E-box only protein 2 (FBXO2), mRNA
VM 033332	Homo sapiens CDC14 cell division cycle 14 homolog B (S. cerevisiae)
VIVI_U33334	(CDC14P) transcript variant 3 mRNA
D.f. 022221	Homo sapiens CDC14 cell division cycle 14 homolog B (S. cerevisiae)
VIM_033331	(CDC14D) transcript variant 2 mKNA
D f 002671	Homo sapiens CDC14 cell division cycle 14 homolog B (S. cerevisiae)
NM_003671	(CDC) 4D) +
7.5.000007	Homo sapiens caspase 4, apoptosis-related cysteine protease (CASP4), transcript
NM_033307	l · 4.1-14- μ-DNΙΛ
	variant delta, mRNA  Homo sapiens caspase 4, apoptosis-related cysteine protease (CASP4), transcript
NM_033306	Homo sapiens caspase 4, apoptosis-iciated dystolic protects
	variant gamma, mRNA  Homo sapiens caspase 4, apoptosis-related cysteine protease (CASP4), transcript
NM_001225	Homo sapiens caspase 4, apoptosis-leiated cysteme protease (S1251-7),
	variant alpha, mRNA
NM_002948	Homo sapiens ribosomal protein L15 (RPL15), mRNA
NM 033228	Homo sapiens ADP-ribosylation factor domain protein 1, 64kD (ARFD1),
_	transcript variant gamma, mRNA
NM_033227	Homo sapiens ADP-ribosylation factor domain protein 1, 64kD (ARFD1),
_	two porint variant heta mRNA
NM_001656	Homo sapiens ADP-ribosylation factor domain protein 1, 64kD (ARFD1),
111.12_001001	transcript variant alpha, mRNA
NM 021203	Home senions APMCE1 protein (APMCE1), mRNA
NM 012095	Homo sapiens adaptor-related protein complex 3, mu 1 subunit (AP3M1),
14141_012000	mRNA
NM 001025	VI-ves serious ribosomal protein \$23 (RP\$23), mRNA
NM 032989	Harma agricing RCI 2-antagonist of cell death (BAD), transcript variant 2, Inches
NM 004322	Homo saniens BCL2-antagonist of cell death (BAD), transcript variant 1, increase
	Homo sapiens death-associated protein kinase 2 (DAPK2), mRNA
NM_014326	Homo sapiens sec22 homolog (SEC22A), mRNA
NM_012430	Homo sapiens sec13-like protein (SEC13L), mRNA
NM_031216	Homo sapiens sec 13-like protein (SEC 13), mRNA  Homo sapiens regulator of G-protein signalling 13 (RGS13), mRNA
NM_002927	Homo sapiens testis expressed sequence 13A (TEX13A), mRNA
NM_031274	Homo sapiens testis expressed sequence 1911 (1911), mRNA
NM_001730	Homo sapiens Kruppel-like factor 5 (intestinal) (KLF5), mRNA  Homo sapiens Kruppel-like factor 5 (intestinal) (KLF5), mRNA
NM_032674	Homo sapiens leucine rich repeat (in FLII) interacting protein 1 (LRRFIP1),
	mRNA
NM_031361	Homo sapiens collagen, type IV, alpha 3 (Goodpasture antigen) binding protein
	(COL4A3BP), transcript variant 2, mRNA
NM 031266	Homo sapiens heterogeneous nuclear ribonucleoprotein A/B (HNRPAB),
_	two page interest 1 mRNA
NM 004499	Homo sapiens heterogeneous nuclear ribonucleoprotein A/B (HNRPAB),
	transcript variant 2 mRNA
NM 004990	Home conjens methionine tRNA synthetase (MARS), mRNA
NM_031244	Homo sapiens sirtuin silent mating type information regulation 2 homolog 5 (3)
14141_021244	
ND4 012241	Homo sapiens sirtuin silent mating type information regulation 2 homolog 5 (S.
NM_012241	gereviciae) (SIRT5) transcript variant 1, mKNA
	The second of th
NM 006845	

NM_030920	Homo sapiens lecuine-rich acidic protein-like protein (LANP-L), mRNA
NM_016228	Homo sapiens L-kynurenine/alpha-aminoadipate aminotransferase (KATII),
<u> </u>	mRNA
NM_017951	Homo sapiens hypothetical protein FLJ20297 (FLJ20297), mRNA
NM_000778	Homo sapiens cytochrome P450, subfamily IVA, polypeptide 11 (CYP4A11),
	mRNA
NM_006582	Homo sapiens glucocorticoid modulatory element binding protein 1 (GMEB1),
	transcript variant 1, mRNA
NM_024482	Homo sapiens glucocorticoid modulatory element binding protein 1 (GMEB1),
37.	transcript variant 2, mRNA
NM_024885	Homo sapiens TAF7-like RNA polymerase II, TATA box binding protein
37.000	(TBP)-associated factor, 50 kD (TAF7L), mRNA
NM_005736	Homo sapiens ARP1 actin-related protein 1 homolog A, centractin alpha (yeast)
77.5 01.100	(ACTR1A), mRNA
NM_014031	Homo sapiens VLCS-H1 protein (VLCS-H1), mRNA
NM_022334	Homo sapiens integrin cytoplasmic domain-associated protein 1 (ICAP-1A),
77.	transcript variant 2, mRNA
NM_007036	Homo sapiens endothelial cell-specific molecule 1 (ESM1), mRNA
NM_006817	Homo sapiens chromosome 12 open reading frame 8 (C12orf8), mRNA
NM_022802	Homo sapiens C-terminal binding protein 2 (CTBP2), transcript variant 2,
37.5 001051	mRNA
NM_001951	Homo sapiens E2F transcription factor 5, p130-binding (E2F5), mRNA
NM_022142	Homo sapiens epididymal sperm binding protein 1 (ELSPBP1), mRNA
NM_012200	Homo sapiens beta-1,3-glucuronyltransferase 3 (glucuronosyltransferase I)
) D ( 000055	(B3GAT3), mRNA
NM_022375	Homo sapiens oculomedin (OCLM), mRNA
NM_004962	Homo sapiens growth differentiation factor 10 (GDF10), mRNA
NM_007372	Homo sapiens RNA helicase-related protein (RNAHP), mRNA
NM_005613	Homo sapiens regulator of G-protein signalling 4 (RGS4), mRNA
NM_006083	Homo sapiens IK cytokine, down-regulator of HLA II (IK), mRNA
NM_012426	Homo sapiens splicing factor 3b, subunit 3, 130kD (SF3B3), mRNA
NM_018164	Homo sapiens hypothetical protein FLJ10637 (FLJ10637), mRNA
NM_006367	Homo sapiens adenylyl cyclase-associated protein (CAP), mRNA
NM_021106	Homo sapiens regulator of G-protein signalling 3 (RGS3), mRNA
NM_021082	Homo sapiens solute carrier family 15 (H+/peptide transporter), member 2
NM_016578	(SLC15A2), mRNA
NM 006671	Homo sapiens HBV pX associated protein-8 (LOC51773), mRNA
14147_0000/1	Homo sapiens solute carrier family 1 (glutamate transporter), member 7 (SLC1A7), mRNA
NM 020650	
NM 015990	Homo sapiens hypothetical protein LOC57333 (LOC57333), mRNA
NM 020905	Homo sapiens lymphocyte activation-associated protein (LOC51088), mRNA Homo sapiens PAN2 protein (PAN2), mRNA
NM_020685	Homo sapiens HT021 (HT021), mRNA
NM_020682	Homo sapiens Cyt19 protein (Cyt19), mRNA
NM 020678	Homo sapiens HT017 protein (HT017), mRNA
NM 020669	
NM 003760	Homo sapiens uncharacterized gastric protein ZA52P (LOC57399), mRNA
14141_002\00	Homo sapiens eukaryotic translation initiation factor 4 gamma, 3 (EIF4G3), mRNA
NM 020412	Homo sapiens CHMP1.5 protein (CHMP1.5), mRNA
NM 020411	Homo sapiens XAGE-1 protein (XAGE-1), mRNA
NM 020408	Homo sapiens CGI-203 protein (CGI-203), mRNA
NM 020395	Homo sapiens hypothetical nuclear factor SBBI22 (LOC57117), mRNA
2.112_020373	Tromo supreme hypothetical flucteal factor SBB122 (LOC5/11/), mKNA

NM 020387	Homo sapiens CATX-8 protein (CATX-8), mRNA
	Homo sapiens cell death regulator aven (LOC57099), mRNA
NM_020371	Homo sapiens HT014 (HT014), mRNA
NM_020362	Homo sapiens cyclin L ania-6a (LOC57018), mRNA
NM_020307	Homo sapiens WW domain binding protein 4 (formin binding protein 21)
NM_007187	(WBP4), mRNA
NM_005644	Homo sapiens TAF12 RNA polymerase II, TATA box binding protein (TBP)-associated factor, 20 kD (TAF12), mRNA
NM_020150	Homo sapiens SAR1 protein (SAR1), mRNA
NM 020167	Homo sapiens neuromedin U receptor 2 (NMU2R), mRNA
NM 020233	Homo sapiens x 006 protein (MDS006), mRNA
NM 020232	Homo sapiens x 003 protein (MDS003), mRNA
NM 020247	Homo sapiens hypothetical protein, clone
_	Telethon(Italy_B41)_Strait02270_FL142 (LOC56997), mRNA
NM_020213	Homo sapiens hypothetical protein from EUROIMAGE 1977056 (LOC56965), mRNA
NM 020153	Homo sapiens hypothetical protein (LOC56912), mRNA
NM_020149	Homo sapiens Meis1, myeloid ecotropic viral integration site 1 homolog 2 (mouse) (MEIS2), mRNA
NM_020120	Homo sapiens UDP-glucose ceramide glucosyltransferase-like 1 (UGCGL1), mRNA
NM_020190	Homo sapiens HNOEL-iso protein (HNOEL-iso), mRNA
NM_020242	Homo sapiens kinesin-like 7 (KNSL7), mRNA
NM_020194	Homo sapiens GL004 protein (GL004), mRNA
NM 020193	Homo sapiens GL002 protein (GL002), mRNA
NM 020189	Homo sapiens DC6 protein (DC6), mRNA
NM 020188	Homo sapiens DC13 protein (DC13), mRNA
NM_020134	Homo sapiens collapsin response mediator protein-5; CRMP3-associated molecule (CRMP5), mRNA
NM 019893	Homo sapiens mitochondrial ceramidase (ASAH2), mRNA
NM 019846	Homo sapiens CC chemokine CCL28 (SCYA28), mRNA
NM 019852	Homo sapiens putative methyltransferase (M6A), mRNA
NM 013338	Homo sapiens Alg5, S. cerevisiae, homolog of (ALG5), mRNA
NM 013341	Homo sapiens hypothetical protein (PTD004), mRNA
NM 013318	Homo sapiens hypothetical protein (LQFBS-1), mRNA
NM 013302	Homo sapiens elongation factor-2 kinase (HSU93850), mRNA
NM_013299	Homo sapiens protein predicted by clone 23627 (HSU79266), mRNA
NM_013347	Homo sapiens replication protein A complex 34 kd subunit homolog Rpa4 (HSU24186), mRNA
NM 019011	Homo sapiens TRIAD3 protein (TRIAD3), mRNA
NM_018965	Homo sapiens triggering receptor expressed on myeloid cells 2 (TREM2), mRNA
NM 019043	Homo sapiens similar to proline-rich protein 48 (LOC54518), mRNA
NM 019006	Homo sapiens protein associated with PRK1 (AWP1), mRNA
NM_019101	Homo sapiens apolipoprotein M (G3A), mRNA
NM 019049	Homo sapiens hypothetical protein (FLJ20054), mRNA
NM 018992	Homo sapiens hypothetical protein (FLJ20040), mRNA
NM 019033	Homo sapiens hypothetical protein (FLJ11235), mRNA
NM 019045	Homo sapiens similar to rab11-binding protein (FLJ11116), mRNA
NM 019079	Homo sapiens hypothetical protein (FLJ10884), mRNA
NM 019073	Homo sapiens hypothetical protein (FLJ10007), mRNA
NM 014298	Homo sapiens quinolinate phosphoribosyltransferase (nicotinate-nucleotide
- TEN	1 A A

(QPCT),
1),
1),
1),
ein
ein
ein
, mRNA
, mRNA
<u> </u>
RNA
),
mRNA
mRNA
12),
- <del>-</del> /,

	DY 120(71 (EI 120(71) mPNA
NM_017924	Homo sapiens hypothetical protein FLJ20671 (FLJ20671), mRNA
NM_017923	Homo sapiens hypothetical protein FLJ20668 (FLJ20668), mRNA
NM_017922	Homo sapiens hypothetical protein FLJ20666 (FLJ20666), mRNA
NM_017908	Homo sapiens hypothetical protein FLJ20626 (FLJ20626), mRNA
NM_017906	Homo sapiens hypothetical protein FLJ20624 (FLJ20624), mRNA
NM_017904	Homo sapiens hypothetical protein FLJ20619 (FLJ20619), mRNA
NM_017890	Homo sapiens hypothetical protein FLJ20583 (FLJ20583), mRNA
NM 017887	Homo sapiens hypothetical protein FLJ20580 (FLJ20580), mRNA
NM_017886	Homo sapiens hypothetical protein FLJ20574 (FLJ20574), mRNA
NM 017880	Homo sapiens hypothetical protein FLJ20558 (FLJ20558), mRNA
NM_017878	Homo sapiens HRAS-like suppressor 2 (HRASLS2), mRNA
NM_017877	Homo sapiens hypothetical protein FLJ20555 (FLJ20555), mRNA
NM_017875	Homo sapiens hypothetical protein FLJ20551 (FLJ20551), mRNA
NM_017870	Homo sapiens hypothetical protein FLJ20539 (FLJ20539), mRNA
NM_017867	Homo sapiens hypothetical protein FLJ20534 (FLJ20534), mRNA
NM_017864	Homo sapiens hypothetical protein FLJ20530 (FLJ20530), mRNA
NM_017857	Homo sapiens slingshot 3 (SSH-3), mRNA
NM_017852	Homo sapiens NALP2 protein (NALP2), mRNA
NM_017850	Homo sapiens hypothetical protein FLJ20508 (FLJ20508), mRNA
NM_017846	Homo sapiens tRNA selenocysteine associated protein (SECP43), mRNA
NM_017841	Homo sapiens hypothetical protein FLJ20487 (FLJ20487), mRNA
NM_017839	Homo sapiens hypothetical protein FLJ20481 (FLJ20481), mRNA
NM_017837	Homo sapiens hypothetical protein FLJ20477 (FLJ20477), mRNA
NM_017832	Homo sapiens hypothetical protein FLJ20457 (FLJ20457), mRNA
NM_017827	Homo sapiens hypothetical protein FLJ20450 (FLJ20450), mRNA
NM_017826	Homo sapiens hypothetical protein FLJ20449 (FLJ20449), mRNA
NM_017823	Homo sapiens hypothetical protein FLJ20442 (FLJ20442), mRNA  Homo sapiens hypothetical protein FLJ20442 (FLJ20442), mRNA
NM_017822	Homo sapiens hypothetical protein FLJ20436 (FLJ20436), mRNA  Homo sapiens hypothetical protein FLJ20436 (FLJ20436), mRNA
NM_017821	Homo sapiens hypothetical protein FLJ20435 (FLJ20435), mRNA  Homo sapiens hypothetical protein FLJ20434 (FLJ20434) mRNA
NM_017815	Homo sapiens hypothetical protein FLJ20424 (FLJ20424), mRNA  Homo sapiens hypothetical protein FLJ20419 (FLJ20419) mRNA
NM_017811	Homo sapiens hypothetical protein FLJ20419 (FLJ20419), mRNA  Homo sapiens hypothetical protein FLJ20417 (FLJ20417) mRNA
NM_017810	Homo sapiens hypothetical protein FLJ20417 (FLJ20417), mRNA  Homo sapiens hypothetical protein FLJ20397 (FLJ20397), mRNA  Homo sapiens hypothetical protein FLJ20397 (FLJ20397), mRNA
NM_017802	Homo sapiens hypothetical protein FLJ20377 (FLJ2037), mRNA  Homo sapiens hypothetical protein FLJ20373 (FLJ20373), mRNA
NM_017792	Homo sapiens regulator of G-protein signalling 3 (RGS3), mRNA
NM_017790	Homo sapiens regulator of G-protein signature 5 (RGSS), in RNA  Homo sapiens hypothetical protein FLJ20366 (FLJ20366), mRNA
NM_017786	Homo sapiens hypothetical protein FLJ20364 (FLJ20364), mRNA  Homo sapiens hypothetical protein FLJ20364 (FLJ20364), mRNA
NM_017785	Homo sapiens hypothetical protein FLJ20343 (FLJ20343), mRNA  Homo sapiens hypothetical protein FLJ20343 (FLJ20343), mRNA
NM_017775	Homo sapiens hypothetical protein FLJ20342 (FLJ20342), mRNA
NM_017774	Homo sapiens hypothetical protein FLJ20337 (FLJ20337), mRNA  Homo sapiens hypothetical protein FLJ20337 (FLJ20337), mRNA
NM_017772	Homo sapiens elongation of very long chain fatty acids (FEN1/Elo2, SUR4/Elo3,
NM_017770	yeast)-like 2 (ELOVL2), mRNA
NR 6 017762	Homo sapiens hypothetical protein FLJ20313 (FLJ20313), mRNA
NM_017762	
NM_017759	
NM_017756	
NM 017753	Homo sapiens hypothetical protein FLJ20297 (FLJ20297), mRNA
NM_017751	
NM 017748	
NM_017744	TT 100070 (TT 100070) mDNA
NM_017740	
NM_017738	Homo sapions hypotherion protein 2 222270 (2 222277)

NB4 017726	TI TO TO TO THE TOTAL OF THE TO
NM_017736	Homo sapiens hypothetical protein FLJ20274 (FLJ20274), mRNA
NM_017735	Homo sapiens hypothetical protein FLJ20272 (FLJ20272), mRNA
NM_017719	Homo sapiens hypothetical protein FLJ20224 (FLJ20224), mRNA
NM_017718	Homo sapiens hypothetical protein FLJ20220 (FLJ20220), mRNA
NM_017716	Homo sapiens membrane-spanning 4-domains, subfamily A, member 12 4-
) D. C. 17711	domains, subfamily A, member 7 (MS4A12), mRNA
NM_017711	Homo sapiens hypothetical protein FLJ20207 (FLJ20207), mRNA
NM_017709	Homo sapiens hypothetical protein FLJ20202 (FLJ20202), mRNA
NM_017704	Homo sapiens hypothetical protein FLJ20189 (FLJ20189), mRNA
NM_017699	Homo sapiens hypothetical protein FLJ20174 (FLJ20174), mRNA
NM_017697	Homo sapiens hypothetical protein FLJ20171 (FLJ20171), mRNA
NM_017687	Homo sapiens hypothetical protein FLJ20147 (FLJ20147), mRNA
NM_017686	Homo sapiens ganglioside induced differentiation associated protein 2 (GDAP2),
NTM 017670	mRNA
NM_017678	Homo sapiens hypothetical protein FLJ20127 (FLJ20127), mRNA
NM_017677	Homo sapiens hypothetical protein FLJ20126 (FLJ20126), mRNA
NM_017676	Homo sapiens hypothetical protein FLJ20125 (FLJ20125), mRNA
NM 017670	Homo sapiens hypothetical protein FLJ20113 (FLJ20113), mRNA
NM_017669	Homo sapiens hypothetical protein FLJ20105 (FLJ20105), mRNA
NM_017665	Homo sapiens hypothetical protein FLJ20094 (FLJ20094), mRNA
NM_017659	Homo sapiens hypothetical protein FLJ20084 (FLJ20084), mRNA
NM_017657	Homo sapiens hypothetical protein FLJ20080 (FLJ20080), mRNA
NM_017645	Homo sapiens hypothetical protein FLJ20060 (FLJ20060), mRNA
NM 017640	Homo sapiens hypothetical protein FLJ20048 (FLJ20048), mRNA
NM_017637	Homo sapiens hypothetical protein FLJ20043 (FLJ20043), mRNA
NM_017636	Homo sapiens transient receptor potential cation channel, subfamily M, member
NM 017634	4 (TRPM4), mRNA
NM 017629	Homo sapiens hypothetical protein FLJ20038 (FLJ20038), mRNA
NM 017622	Homo sapiens hypothetical protein FLJ20033 (FLJ20033), mRNA
NM 017620	Homo sapiens hypothetical protein FLJ20014 (FLJ20014), mRNA Homo sapiens hypothetical protein FLJ20011 (FLJ20011), mRNA
NM_018396	Homo sapiens putative methyltransferase (METL), mRNA
NM 018381	Homo sapiens hypothetical protein FLJ11286 (FLJ11286), mRNA
NM_018371	Homo sapiens hypothetical protein FLJ11260 (FLJ11260), mRNA  Homo sapiens hypothetical protein FLJ11264 (FLJ11264), mRNA
NM_018368	Homo sapiens hypothetical protein FLJ11240 (FLJ11240), mRNA  Homo sapiens hypothetical protein FLJ11240 (FLJ11240), mRNA
NM 018367	Homo sapiens phytoceramidase, alkaline (PHCA), mRNA
NM 018364	Homo sapiens hypothetical protein FLJ11220 (FLJ11220), mRNA
NM 018363	Homo sapiens hypothetical protein FLJ11220 (FLJ11220), mRNA  Homo sapiens hypothetical protein FLJ11218 (FLJ11218), mRNA
NM 018361	Homo sapiens hypothetical protein FLJ11210 (FLJ11210), mRNA  Homo sapiens hypothetical protein FLJ11210 (FLJ11210), mRNA
NM_018358	Homo sapiens hypothetical protein FLJ11210 (FLJ11210), mRNA  Homo sapiens hypothetical protein FLJ11198 (FLJ11198), mRNA
NM_018353	Homo sapiens hypothetical protein FLJ11186 (FLJ11186), mRNA
NM 018352	Homo sapiens hypothetical protein FLJ11186 (FLJ11186), mRNA  Homo sapiens hypothetical protein FLJ11184 (FLJ11184), mRNA
NM 018340	Homo sapiens hypothetical protein FLJ11151 (FLJ11151), mRNA
NM 018339	Homo sapiens hypothetical protein FLJ11151 (FLJ11151), mRNA  Homo sapiens hypothetical protein FLJ11149 (FLJ11149), mRNA
NM 018336	Homo sapiens hypothetical protein FLJ11136 (FLJ11136), mRNA
NM 018333	Homo sapiens hypothetical protein FLJ11136 (FLJ11136), mRNA  Homo sapiens hypothetical protein FLJ20666 (FLJ20666), mRNA
NM_018332	Homo sapiens hypothetical protein ET 111126 (ET 11126), mRNA
NM_018330	Homo sapiens hypothetical protein FLJ11126 (FLJ11126), mRNA Homo sapiens KIAA1598 protein (KIAA1598), mRNA
NM_018322	Homo saniens hymothetical protein EX 11101 (EX 11101)
NM_018318	Homo sapiens hypothetical protein FLJ11101 (FLJ11101), mRNA
NM_018310	Homo sapiens hypothetical protein FLJ11088 (FLJ11088), mRNA
7.11.1 O 100 10	Homo sapiens BRF2, subunit of RNA polymerase III transcription initiation factor, BRF1-like (BRF2), mRNA
L	THEOLI, DIG 1-TIRE (DRF2), IIIKNA

NM_018303	Homo sapiens hypothetical protein FLJ11026 (FLJ11026), mRNA
NM_018298	Homo sapiens hypothetical protein FLJ11006 (FLJ11006), mRNA
NM_018287	Homo sapiens hypothetical protein FLJ10971 (FLJ10971), mRNA
NM_018286	Homo sapiens hypothetical protein FLJ10970 (FLJ10970), mRNA
NM_018283	Homo sapiens hypothetical protein FLJ10956 (FLJ10956), mRNA
NM_018281	Homo sapiens hypothetical protein FLJ10948 (FLJ10948), mRNA
NM_018278	Homo sapiens hypothetical protein FLJ10933 (FLJ10933), mRNA
NM_018276	Homo sapiens slingshot 3 (SSH-3), mRNA
NM_018273	Homo sapiens hypothetical protein FLJ10922 (FLJ10922), mRNA
NM_018272	Homo sapiens hypothetical protein FLJ10921 (FLJ10921), mRNA
NM_018268	Homo sapiens hypothetical protein FLJ10904 (FLJ10904), mRNA
NM_018265	Homo sapiens hypothetical protein FLJ10901 (FLJ10901), mRNA
NM_018254	Homo sapiens hypothetical protein FLJ10876 (FLJ10876), mRNA
NM_018253	Homo sapiens hypothetical protein FLJ10875 (FLJ10875), mRNA
NM_018252	Homo sapiens hypothetical protein FLJ10874 (FLJ10874), mRNA
NM_018245	Homo sapiens hypothetical protein FLJ10851 (FLJ10851), mRNA
NM_018241	Homo sapiens hypothetical protein FLJ10846 (FLJ10846), mRNA
NM_018239	Homo sapiens hypothetical protein FLJ10751 (FLJ10751), mRNA
NM_018230	Homo sapiens nucleoporin 133kD (NUP133), mRNA
NM_018223	Homo sapiens checkpoint with forkhead and ring finger domains (CHFR),
	mRNA
NM_018219	Homo sapiens hypothetical protein FLJ10786 (FLJ10786), mRNA
NM_018217	Homo sapiens chromosome 20 open reading frame 31 (C20orf31), mRNA
NM_018212	Homo sapiens likely ortholog of mouse NPC derived proline rich protein 1
ND 6 010011	(FLJ10773), mRNA
NM_018211	Homo sapiens hypothetical protein FLJ10770 (KIAA1579), mRNA
NM_018207	Homo sapiens hypothetical protein FLJ10759 (FLJ10759), mRNA
NM_018205	Homo sapiens hypothetical protein FLJ10751 (FLJ10751), mRNA Homo sapiens hypothetical protein FLJ10718 (FLJ10718), mRNA
NM_018192	Homo sapiens hypothetical protein FLJ10718 (FLJ10718), mRNA  Homo sapiens hypothetical protein FLJ10709 (FLJ10709), mRNA
NM_018188 NM_018187	Homo sapiens hypothetical protein FLJ10709 (FLJ10709), mRNA  Homo sapiens hypothetical protein FLJ10707 (FLJ10707), mRNA
NM_018186	Homo sapiens hypothetical protein FLJ10707 (FLJ10707), mRNA  Homo sapiens hypothetical protein FLJ10706 (FLJ10706), mRNA
NM 018184	Homo sapiens hypothetical protein FLJ10702 (FLJ10702), mRNA
NM 018179	Homo sapiens hypothetical protein FLJ10688 (FLJ10688), mRNA
NM_018179	Homo sapiens hypothetical protein FLJ10687 (FLJ10687), mRNA
NM 018169	Homo sapiens hypothetical protein FLJ10657 (FLJ10657), mRNA
NM 018161	Homo sapiens hypothetical protein FLJ10631 (FLJ10631), mRNA
NM 018159	Homo sapiens hypothetical protein FLJ10628 (FLJ10628), mRNA
NM 018147	Homo sapiens hypothetical protein FLJ10582 (FLJ10582), mRNA
NM_018142	Homo sapiens hypothetical protein FLJ10569 (FLJ10569), mRNA
NM 018137	Homo sapiens protein arginine N-methyltransferase 6 (PRMT6), mRNA
NM_018136	Homo sapiens hypothetical protein FLJ10517 (FLJ10517), mRNA
NM 018133	Homo sapiens hypothetical protein FLJ10546 (FLJ10546), mRNA
NM 018122	Homo sapiens hypothetical protein FLJ10514 (FLJ10514), mRNA
NM 018120	Homo sapiens hypothetical protein FLJ10511 (FLJ10511), mRNA
NM 018119	Homo sapiens hypothetical protein FLJ10509 (FLJ10509), mRNA
NM 018116	Homo sapiens misato (FLJ10504), mRNA
NM 018112	Homo sapiens hypothetical protein FLJ10493 (FLJ10493), mRNA
NM_018106	Homo sapiens hypothetical protein FLJ10479 (FLJ10479), mRNA
NM 018101	Homo sapiens hypothetical protein FLJ10468 (FLJ10468), mRNA
NM 018100	Homo sapiens hypothetical protein FLJ10466 (FLJ10466), mRNA
NM 018099	Homo sapiens hypothetical protein FLJ10462 (FLJ10462), mRNA
	The second of th

NB4 010007	
NM_018097	Homo sapiens hypothetical protein FLJ10460 (FLJ10460), mRNA
NM 018093	Homo sapiens hypothetical protein FLJ10439 (FLJ10439), mRNA
NM_018092	Homo sapiens hypothetical protein FLJ10430 (FLJ10430), mRNA
NM_018091	Homo sapiens hypothetical protein FLJ10422 (FLJ10422), mRNA
NM_018090	Homo sapiens hypothetical protein FLJ10420 (FLJ10420), mRNA
NM_018087	Homo sapiens hypothetical protein FLJ10407 (FLJ10407), mRNA
NM_018086	Homo sapiens fidgetin (FIGN), mRNA
NM_018078	Homo sapiens hypothetical protein FLJ10378 (FLJ10378), mRNA
NM_018076	Homo sapiens hypothetical protein FLJ10376 (FLJ10376), mRNA
NM_018075	Homo sapiens hypothetical protein FLJ10375 (FLJ10375), mRNA
NM_018072	Homo sapiens hypothetical protein FLJ10359 (FLJ10359), mRNA
NM_018070	Homo sapiens hypothetical protein FLJ10355 (FLJ10355), mRNA
NM_018060	Homo sapiens hypothetical protein FLJ10326 (FLJ10326), mRNA
NM_018054	Homo sapiens homolog of rat nadrin (RICH1), mRNA
NM_018052	Homo sapiens hypothetical protein FLJ10305 (FLJ10305), mRNA
NM_018051	Homo sapiens hypothetical protein FLJ10300 (FLJ10300), mRNA
NM_018047	Homo sapiens hypothetical protein FLJ10290 (FLJ10290), mRNA
NM_018043	Homo sapiens hypothetical protein FLJ10261 (FLJ10261), mRNA
NM_018040	Homo sapiens hypothetical protein FLJ10252 (FLJ10252), mRNA
NM_018039	Homo sapiens hypothetical protein FLJ10251 (FLJ10251), mRNA
NM_018038	Homo sapiens hypothetical protein FLJ10246 (FLJ10246), mRNA
NM_018035	Homo sapiens hypothetical protein FLJ10241 (FLJ10241), mRNA
NM_018034	Homo sapiens hypothetical protein FLJ10233 (FLJ10233), mRNA
NM_018033	Homo sapiens hypothetical protein FLJ10232 (FLJ10232), mRNA
NM_018026	Homo sapiens hypothetical protein FLJ10209 (FLJ10209), mRNA
NM_018025	Homo sapiens hypothetical protein FLJ10206 (FLJ10206), mRNA
NM_018011	Homo sapiens hypothetical protein FLJ10154 (FLJ10154), mRNA
NM_018009	Homo sapiens hypothetical protein FLJ10143 (FLJ10143), mRNA
NM_018008	Homo sapiens hypothetical protein FLJ10142 (FLJ10142), mRNA
NM_018001	Homo sapiens hypothetical protein FLJ10120 (FLJ10120), mRNA
NM_017994	Homo sapiens hypothetical protein FLJ10099 (FLJ10099), mRNA
NM_017993	Homo sapiens hypothetical protein FLJ10094 (FLJ10094), mRNA
NM_017988	Homo sapiens hypothetical protein FLJ10074 (FLJ10074), mRNA
NM_017987	Homo sapiens Run- and FYVE-domain containing protein (Rabip4R), mRNA
NM_017976	Homo sapiens hypothetical protein FLJ10038 (FLJ10038), mRNA
NM_018409	Homo sapiens hypothetical protein DKFZp761O0113 (DKFZp761O0113),
ND4 017601	mkna
NM_017601	Homo sapiens hypothetical protein DKFZp761H221 (DKFZp761H221), mRNA
NM_018713	Homo sapiens hypothetical protein DKFZp547M236 (DKFZp547M236), mRNA
NM_017606	Homo sapiens hypothetical protein DKFZp434K1210 (DKFZp434K1210)
ND4 017546	mRNA
NM_017546	Homo sapiens hypothetical protein (C40), mRNA
NM_018458	Homo sapiens uncharacterized bone marrow protein BM042 (BM042), mRNA
NM_018456	Homo sapiens uncharacterized bone marrow protein BM040 (BM040), mRNA
NM_018455	Homo sapiens uncharacterized bone marrow protein BM039 (BM039), mRNA
NM_018453	Homo sapiens uncharacterized bone marrow protein BM036 (BM036), mRNA
NM_018452	Homo sapiens chromosome 6 open reading frame 35 (C6orf35), mRNA
NM_018489	Homo sapiens hypothetical protein ASH1 (ASH1), mRNA
NM_004227	Homo sapiens pleckstrin homology, Sec7 and coiled/coil domains 3 (PSCD3)
	mrna
NM_007014	Homo sapiens Nedd-4-like ubiquitin-protein ligase (WWP2), mRNA
NM_017431	Homo sapiens protein kinase, AMP-activated, gamma 3 non-catalytic subunit

	(PRKAG3), mRNA
NM 017426	Homo sapiens nucleoporin 54kD (NUP54), mRNA
7.6.01.6050	Homo sapiens testican 3 (HSAJ1454), mRNA
	Homo sapiens methyltransferase COQ3 (COQ3), mRNA  Homo sapiens methyltransferase COQ3 (COQ3), mRNA
NM 006854	Homo sapiens MDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein
_	retention receptor 2 (KDELR2), mRNA
NM_015976	Homo sapiens sorting nexin 7 (SNX7), mRNA  Homo sapiens sorting nexin 7 (SNX7), mRNA
NM 016577	Homo saniens RAB6B, member RAS oncogene railing (RAB6B), mid 41
NM 016559	Uomo saniens PXR2h protein (PXR2b), mRNA
NM 016297	Homo sapiens prenylcysteine lyase (PCL1), mRNA
NM 016524	R/V protein (1.0(5)/60), $R/V$
NM 016507	Homo saniens CDC2-related protein kinase / (CIKRS), mid VI
NM 016446	Harris genians NAG-5 protein (LOC51/54), mRNA
NM 016382	· · · · · · · · · · · · · · · · · · ·
NM_016354	Homo sapiens solute carrier family 21 (organic anion transporter), member 12
1414101035 .	(GY CO1 4 12) PN 4
NM 016298	diagona related protein (LOC) L/20), IRRIVA
NM 016290	Transiens retinged v recentor interacting Dioletti (LOC51720), indiaz
NM 016280	t == the confloaterace related protein (LUC)1/10), IIIXVI
NM 016229	to a second prome by reductase by R. 2 (LUC) 1/00), 11101111
NM 016213	The state of the s
	Try and suppressor of fused homolog (Diosophila) (5010), hitch
NM_016169 NM_016084	Homo saniens RAS dexamethasone-induced I (RASDI), INCVI
	Homo saniens CGI-147 protein (LOC51651), mRNA
NM_016077	T
NM_016023	Theme geniens non-canonical ubquitin conjugating enzyme i (1000001); indicating
NM_016021	TOTATION 11 54 profess (1)K P/P434J134J, HUMAN
NM_016003	Homo sapiens DKFZF434J134 protein (Sta 22) Homo sapiens calcium/calmodulin-dependent protein kinase (CaM kinase) II
NM_015981	alpha (CAMK2A) mRNA
27.5.015040	T CCI 20 protein (1 OC > 16UX) MKNA
NM_015949	Homo coniens dickkont homolog 3 (Aenopus laevis) (DRRS); me u
NM_015881	Homo sapiens hypothetical protein (LOC51316), mRNA
NM_016619	Homo sapiens DHHC1 protein (LOC51304), mRNA
NM_016598	Homo sapiens M5-14 protein (LOC51300), mRNA
NM_016589	Homo sapiens neuritin (LOC51299), mRNA
NM_016588	Homo sapiens neutrin (EGSTES), mRNA  Homo sapiens peptide transporter 3 (PHT2), mRNA
NM_016582	Homo sapiens CDA14 (LOC51290), mRNA
NM_016570	Homo sapiens CDA14 (EOCS1220), mRNA  Homo sapiens E2IG2 protein (LOCS1287), mRNA
NM_016565	Homo sapiens apoptosis regulator (LOC51283), mRNA
NM_016561	Homo sapiens apoptosis regulator (EGCs 1200), mRNA
NM_016526	Homo sapiens GS15 (LOC51272), mRNA  Homo sapiens pipecolic acid oxidase (PIPOX), mRNA  Homo sapiens pipecolic acid oxidase (PIPOX), mRNA
NM_016518	Homo sapiens pipecone acid oxidase (Tit 622);
NM_016495	Homo sapiens hypothetical protein (LOC51256), mRNA  Homo sapiens hypothetical protein (LOC51249), mRNA
NM_016486	
NM_016477	
NM_016465	
NM_016456	
NM_016350	
NM 016274	Homo saniens CK2 interacting protein 1; HQ0024c protein (LOCS1177), max-
NM 016261	
NM 016216	Homo saniens debranching enzyme homolog I (S. cerevisiae) (DDRI), mid 12
NM 016208	$\frac{1}{2}$
NM 016206	

NM_016185	Homo sapiens hematological and neurological expressed 1 (HN1), mRNA
NM_016181	Homo sapiens melanoma antigen (LOC51152), mRNA
NM_016139	Homo sapiens 16.7Kd protein (LOC51142), mRNA
NM_016129	Homo sapiens COP9 constitutive photomorphogenic homolog subunit 4
	(Arabidopsis) (COPS4), mRNA
NM_016122	Homo sapiens NY-REN-58 antigen (LOC51134), mRNA
NM_016119	Homo sapiens putative zinc finger protein NY-REN-34 antigen (LOC51131),
	mRNA
NM_016103	Homo sapiens GTP-binding protein Sara (LOC51128), mRNA
NM_016099	Homo sapiens HSPC041 protein (LOC51125), mRNA
NM_016096	Homo sapiens HSPC038 protein (LOC51123), mRNA
NM_016037	Homo sapiens CGI-94 protein (LOC51118), mRNA
NM_016014	Homo sapiens CGI-67 protein (LOC51104), mRNA
NM_015997	Homo sapiens CGI-41 protein (LOC51093), mRNA
NM 015974	Homo sapiens lambda-crystallin (LOC51084), mRNA
NM_015973	Homo sapiens galanin-related peptide (LOC51083), mRNA
NM_015972	Homo sapiens RNA polymerase I 16 kDa subunit (LOC51082), mRNA
NM_015953	Homo sapiens eNOS interacting protein (NOSIP), mRNA
NM_015936	Homo sapiens CGI-04 protein (LOC51067), mRNA
NM 015895	Homo sapiens geminin (LOC51053), mRNA
NM 015882	Homo sapiens RIG-like 5-6 (LOC51048), mRNA
NM 015853	Homo sapiens ORF (LOC51035), mRNA
NM 016080	Homo sapiens CGI-150 protein (LOC51031), mRNA
NM 016078	Homo sapiens CGI-148 protein (LOC51030), mRNA
NM 016076	Homo sapiens CGI-146 protein (LOC51029), mRNA
NM 016052	Homo sapiens CGI-115 protein (LOC51018), mRNA
NM 016049	Homo sapiens CGI-112 protein (LOC51016), mRNA
NM 015940	Homo sapiens CGI-10 protein (LOC51004), mRNA
NM 016505	Homo sapiens hypothetical protein (HSPC251), mRNA
NM 016485	Homo sapiens hypothetical protein (HSPC228), mRNA
NM 016472	Homo sapiens hypothetical protein (HSPC210), mRNA
NM 016464	Homo sapiens hypothetical protein (HSPC196), mRNA
NM 016462	Homo sapiens hypothetical protein (HSPC194), mRNA
NM 016535	Homo sapiens HSPC189 protein (HSPC189), mRNA
NM 016404	Homo sapiens hypothetical protein (HSPC152), mRNA
NM 016403	Homo sapiens hypothetical protein (HSPC148), mRNA
NM 016399	Homo sapiens hypothetical protein (HSPC132), mRNA
NM_016395	Homo sapiens butyrate-induced transcript 1 (HSPC121), mRNA
NM 016387	Homo sapiens hypothetical protein (HSPC060), mRNA
NM_016101	Homo sapiens hypothetical protein (HSPC031), mRNA
NM 015918	Homo sapiens homolog of yeast RNase MRP/RNase P protein Pop5 (POP5),
1111_013910	mRNA
NM 016257	Homo sapiens hippocalcin-like protein 4 (HPCAL4), mRNA
NM_016287	Homo sapiens HP1-BP74 (HP1-BP74), mRNA
NM_015888	Homo sapiens hook1 protein (HOOK1), mRNA
NM 015852	Homo sapiens Krueppel-related zinc finger protein (H-plk), mRNA
NM_016451	Homo sapiens coatomer protein complex, subunit beta (COPB), mRNA
NM 015986	Homo sapiens cytokine receptor-like factor 3 (CRLF3), mRNA
NM_016204	
	Homo sapiens growth differentiation factor 2 (GDF2), mRNA
NM_016617 NM_014822	Homo sapiens hypothetical protein (BM-002), mRNA
171VI_U14822	Homo sapiens SEC24 related gene family, member D (S. cerevisiae) (SEC24D),
L	mRNA

	TO COOK DATA
NM 014059	Homo sapiens RGC32 protein (RGC32), mRNA
NM 014040	Homo sapiens PTD015 protein (PTD015), mRNA
NM 014039	Homo sapiens PTD012 protein (PTD012), mRNA
NM 014111	Homo sapiens PRO2086 protein (PRO2086), mRNA
NM 014106	Homo sapiens PRO1914 protein (PRO1914), mRNA
NM 014104	Homo sapiens PRO1880 protein (PRO1880), mRNA
NM 014100	Homo sapiens PRO1770 protein (PRO1770), mRNA
NM 014137	Homo sapiens PRO0650 protein (PRO0650), mRNA
NM_014127	Homo sapiens PRO0456 protein (PRO0456), mRNA
NM 014123	Homo sapiens PRO0246 protein (PRO0246), mRNA
NM 014114	Homo sapiens PRO0097 protein (PRO0097), mRNA
NM 014113	Homo sapiens PRO0038 protein (PRO0038), mRNA
NM 014048	Homo sapiens KIAA1243 protein (KIAA1243), mRNA
NM 015368	Homo sapiens pannexin 1 (PANX1), mRNA
NM 014910	Homo sapiens KIAA1084 protein (KIAA1084), mRNA
NM 014916	Homo sapiens KIAA1079 protein (KIAA1079), mRNA
NM 014967	Trans conjens KIA A1018 protein (KIAA1018), mRNA
NM 014953	Homo sapiens mitotic control protein dis3 homolog (KIAA1008), mKNA
NM 014954	Homo sapiens KIAA0985 protein (KIAA0985), mRNA
NM 014917	Homo saniens netrin G1 (KIAA0976), mRNA
NM 014930	Homo sapiens KIAA0972 protein (KIAA0972), mRNA
NM 014907	Homo sapiens KIAA0967 protein (KIAA0967), mRNA
NM 014912	Homo sapiens KIAA0940 protein (KIAA0940), mRNA
NM 014021	Homo sapiens KIAA0923 protein (KIAA0923), mRNA
NM_014899	Homo sapiens KIAA0878 protein (KIAA0878), mRNA
NM 014951	Homo saniens KIAA0844 protein (KIAA0844), mRNA
NM 014729	Homo sapiens KIAA0808 gene product (KIAA0808), mRNA
NM 014813	Homo sapiens KIAA0806 gene product (KIAA0806), mRNA
NM 014829	Homo saniens RNA helicase (KIAA0801), mRNA
NM 014698	Homo sapiens KIAA0792 gene product (KIAA0792), mRNA
NM 014824	Homo sapiens KIAA0769 gene product (KIAA0769), mkNA
NM 014677	Homo sapiens KIAA0751 gene product (KIAA0751), mkNA
NM 014705	Homo sapiens KIAA0716 gene product (KIAA0716), mRNA
NM 014861	Homo sapiens KIAA0703 gene product (KIAA0703), mRNA
NM 014721	Homo saniens KIA A0680 gene product (KIAA0680), mRNA
NM_014827	Homo sapiens KIAA0663 gene product (KIAA0663), mRNA
NM_014645	Homo saniens KIAA0635 gene product (KIAA0635), mkNA
NM 014664	Homo saniens KIAA0615 gene product (KIAA0615), mRNA
NM 014834	Homo saniens KIAA0563 gene product (KIAA0563), mRNA
NM 014696	Homo sapiens KIAA0514 gene product (KIAA0514), mRNA
NM 014732	Homo sapiens KIA A0513 gene product (KIA A0513), mRNA
NM_014710	Homo saniens KIA A0443 gene product (KIAA0443), mRNA
NM_014797	Homo sapiens KIAA0441 gene product (KIAA0441), mRNA
NM_014819	Homo saniens KIAA0438 gene product (KIAA0438), mRNA
NM_015216	Homo sapiens KIAA0433 protein (KIAA0433), mRNA
NM 015251	TY agricus VIA A0431 protein (KIA A0431), mRNA
NM 015185	
14141_012102	mPNIA
NM_014711	Homo saniens KIAA0419 gene product (KIAA0419), mRNA
NM 015564	Homo saniens KIA A0416 protein (KIAA0416), mRNA
NM 014778	
14171 014770	

NM_014659	Homo sapiens KIAA0377 gene product (KIAA0377), mRNA
NM_014639	Homo sapiens KIAA0372 gene product (KIAA0372), mRNA
NM_014786	Homo sapiens KIAA0337 gene product (KIAA0337), mRNA
NM_014845	Homo sapiens KIAA0274 gene product (KIAA0274), mRNA
NM_014745	Homo sapiens KIAA0233 gene product (KIAA0233), mRNA
NM 014643	Homo sapiens KIAA0222 gene product (KIAA0222), mRNA
NM 014674	Homo sapiens KIAA0212 gene product (KIAA0212), mRNA
NM 014720	Homo sapiens Ste20-related serine/threonine kinase (SLK), mRNA
NM 014761	Homo sapiens KIAA0174 gene product (KIAA0174), mRNA
NM 014730	Homo sapiens KIAA0152 gene product (KIAA0152), mRNA
NM 014661	Homo sapiens KIAA0140 gene product (KIAA0140), mRNA
NM 014777	Homo sapiens KIAA0133 gene product (KIAA0133), mRNA
NM 014815	Homo sapiens KIAA0130 gene product (KIAA0130), mRNA
NM 014755	Homo sapiens transcriptional regulator interacting with the PHS-bromodomain 2
	(TRIP-Br2), mRNA
NM_014628	Homo sapiens gene predicted from cDNA with a complete coding sequence
_	(KIAA0110), mRNA
NM 014814	Homo sapiens KIAA0107 gene product (KIAA0107), mRNA
NM 014752	Homo sapiens KIAA0102 gene product (KIAA0102), mRNA
NM 014780	Homo sapiens KIAA0076 gene product (KIAA0076), mRNA
NM 014882	Homo sapiens KIAA0053 gene product (KIAA0053), mRNA
NM 014750	Homo sapiens KIAA0008 gene product (KIAA0008), mRNA
NM 015684	Homo sapiens mitochondrial ATP synthase regulatory component factor B
	(ATPW), mRNA
NM 014186	Homo sapiens HSPC166 protein (HSPC166), mRNA
NM 014184	Homo sapiens HSPC163 protein (HSPC163), mRNA
NM 014181	Homo sapiens HSPC159 protein (HSPC159), mRNA
NM 014179	Homo sapiens HSPC157 protein (HSPC157), mRNA
NM 014166	Homo sapiens HSPC126 protein (HSPC126), mRNA
NM 014155	Homo sapiens HSPC063 protein (HSPC063), mRNA
NM 014038	Homo sapiens HSPC028 protein (HSPC028), mRNA
NM 014017	Homo sapiens HSPC003 protein (HSPC003), mRNA
NM 014053	Homo sapiens FLVCR protein (FLVCR), mRNA
NM 015400	Homo sapiens DKFZP586N0721 protein (DKFZP586N0721), mRNA
NM 015583	Homo sapiens DKFZP586M0622 protein (DKFZP586M0622), mRNA
NM 015485	Homo sapiens DKFZP566K023 protein (DKFZP566K023), mRNA
NM 014043	Homo sapiens DKFZP564O123 protein (DKFZP564O123), mRNA
NM 015387	Homo sapiens preimplantation protein 3 (PREI3), mRNA
NM 014056	Homo sapiens DKFZP564K247 protein (DKFZP564K247), mRNA
NM 015623	Homo sapiens putative ankyrin-repeat containing protein (DKFZP564D166),
14141_015025	mRNA
NM 015582	Homo sapiens DKFZP564B147 protein (DKFZP564B147), mRNA
NM 015610	Homo sapiens DKFZP434J154 protein (DKFZP434J154), mRNA
NM 015590	Homo sapiens DKFZP434F1735 protein (DKFZP434F1735), mRNA
NM 015644	Homo sapiens DKFZP434B103 protein (DKFZP434B103), mRNA
NM 015396	Homo sapiens DKFZP434A043 protein (DKFZP434A043), mRNA
NM 014058	Homo sapiens DESC1 protein (DESC1), mRNA
NM 015680	Homo sapiens bescriptotein (DESCr), mRNA  Homo sapiens hypothetical protein (CGI-57), mRNA
	Homo sapiens brain protein I3 (BRI3), mRNA
NM_015379	Homo sapiens solute carrier family 2, (facilitated glucose transporter) member 8
NM_014580	(SLC2A8), mRNA
NM 014280	Homo sapiens DnaJ (Hsp40) homolog, subfamily C, member 8 (DNAJC8),
11111 01 1200	Charles (110) 10/ nomeroes, exercising 5, memory 6 (111/11/11/11/11/11/11/11/11/11/11/11/11

	mRNA
T 6 01 4212	Hame seniens small membrane protein 1 (SMP1), mRNA
NM 014229	Homo sapiens solute carrier family 6 (neurotransmitter transporter, GABA),
\	momber 11 (SI C6A11) mRNA
T 5 01 4575	Home seniens schwannomin interacting protein 1 (SCHIP1), mRNA
NM 014402	Homo sapiens low molecular mass ubiquinone-binding protein (9.5kD) (QP-C),
	DNIA
NM_014394	Homo sapiens growth hormone inducible transmembrane protein (GHITM),
	DNIA
NM 014225	Homo sapiens protein phosphatase 2 (formerly 2A), regulatory subunit A (PR
	65), alpha isoform (PPP2R1A), mRNA
NM 014497	Homo saniens nuclear protein (NP220), mRNA
NM 014399	Homo saniens tetraspan NET-6 protein (NET-6), mRNA
NM 014889	Homo saniens metalloprotease 1 (pitrilysin family) (MPI), mRNA
NM 014484	Homo sapiens molybdenum cofactor synthesis 3 (MOCS3), mRNA
NM 014447	Homo sapiens arfaptin 1 (HSU52521), mRNA
NM 014350	Homo seniens TNF-induced protein (GG2-1), mRNA
NM 014478	Homo sapiens calcitonin gene-related peptide-receptor component protein
11112_01	(CCDP_PCP) mRNA
NM 014482	Homo sapiers hone morphogenetic protein 10 (BMP10), mRNA
NM 014474	Homo sapiens acid sphingomyelinase-like phosphodiesterase (ASML3B),
11112_02	mRNA
NM 014480	Homo sapiens zinc finger protein (AF020591), mRNA
NM 014576	Homo sapiens Apobec-1 complementation factor; APOBEC-1 stimulating
	protein (ACF), mRNA
NM 005884	Home conjugg n21 (CDK N1 A)-activated kinase 4 (PAK4), mRNA
NM 013434	Homo sapiens calsenilin, presenilin binding protein, EF hand transcription factor
_	(CSFN) mRNA
NM 012446	Homo sapiens single-stranded DNA binding protein 2 (SSBP2), mRNA
NM 013235	Homo saniens putative ribonuclease III (RNASE3L), mRNA
NM 013349	Homo sapiens secreted protein of unknown function (SPUF), mRNA
NM 013323	Homo saniens sorting nexin 11 (SNX11), mRNA
NM 013388	Homo saniens prolactin regulatory element binding (PREB), mRNA
NM 013328	Homo saniens pyrroline 5-carboxylate reductase isotorm (P3CR2), mRNA
NM 013370	Homo saniens pregnancy-induced growth inhibitor (OKL38), mRNA
NM 013277	Home capiens Rac GTPase activating protein 1 (RACGAP1), micha
NM 013285	Homo sapiens nucleolar GTPase (HUMAUANTIG), mRNA
NM 013320	Homo saniens host cell factor 2 (HCF-2), mRNA
NM 013391	Homo saniens dimethylglycine dehydrogenase precursor (DMGDH), mRNA
NM 013253	Homo saniens dickkonf homolog 3 (Xenopus laevis) (DKK3), mRNA
NM 013339	Homo sapiens dolichyl-P-Glc:Man9GlcNAc2-PP-dolichylglucosyltransferase
11117_01000	(ALC6) mRNA
NM 004120	Homo sapiens guanylate binding protein 2, interferon-inducible (GBP2), mRN2
NM 005690	Homo sapiens dynamin 1-like (DNM1L), transcript variant 3, mkNA
NM 012063	Homo sapiens dynamin 1-like (DNM1L), transcript variant 2, mRNA
NM 012470	Homo sapiens transportin-SR (TRN-SR), mRNA
NM 012252	Homo saniens transcription factor EC (TFEC), mRNA
NM_012250	Homo sapiens related RAS viral (r-ras) oncogene homolog 2 (RRAS2), mRNA
NM 012249	Homo sapiens ras-like protein (TC10), mRNA
NM 012249 NM 012388	Homo saniens pallidin homolog (mouse) (PLDN), mRNA
	TIVITO OUPTOIN PULLANTI INCANOLOS (
NM 012322	Homo sapiens U6 snRNA-associated Sm-like protein (LSM5), mRNA Homo sapiens karyopherin alpha 6 (importin alpha 7) (KPNA6), mRNA

ND ( 010100	II conient Chronel orthin II (ECD 2) mPNA
NM_012189	Homo sapiens fibrousheathin II (FSP-2), mRNA
NM_012081	Homo sapiens ELL-RELATED RNA POLYMERASE II, ELONGATION
	FACTOR (ELL2), mRNA
NM_003996	Homo sapiens glutathione peroxidase 5 (epididymal androgen-related protein)
	(GPX5), transcript variant 2, mRNA
NM_005260	Homo sapiens growth differentiation factor 9 (GDF9), mRNA
NM_007352	Homo sapiens elastase 3B, pancreatic (ELA3B), mRNA
NM_006685	Homo sapiens proline rich 3 (PROL3), mRNA
NM_007357	Homo sapiens low density lipoprotein receptor defect C complementing (LDLC), mRNA
NM 004133	Homo sapiens hepatocyte nuclear factor 4, gamma (HNF4G), mRNA
NM_003144	Homo sapiens signal sequence receptor, alpha (translocon-associated protein alpha) (SSR1), mRNA
NM 007324	Homo sapiens MAD, mothers against decapentaplegic homolog (Drosophila)
14141_007524	interacting protein, receptor activation anchor (MADHIP), transcript variant 1, mRNA
NM_007323	Homo sapiens MAD, mothers against decapentaplegic homolog (Drosophila) interacting protein, receptor activation anchor (MADHIP), transcript variant 2,
	mRNA
NM_005162	Homo sapiens angiotensin receptor-like 2 (AGTRL2), mRNA
NM_005501	Homo sapiens integrin, alpha 3 (antigen CD49C, alpha 3 subunit of VLA-3
	receptor) (ITGA3), transcript variant b, mRNA
NM_007144	Homo sapiens zinc finger protein 144 (Mel-18) (ZNF144), mRNA
NM 007286	Homo sapiens synaptopodin (KIAA1029), mRNA
NM 007199	Homo sapiens interleukin-1 receptor-associated kinase M (IRAK-M), mRNA
NM 007283	Homo sapiens monoglyceride lipase (MGLL), mRNA
NM 007241	Homo sapiens EAP30 subunit of ELL complex (EAP30), mRNA
NM_007212	Homo sapiens ring finger protein 2 (RNF2), mRNA
NM 007236	Homo sapiens calcium binding protein P22 (CHP), mRNA
NM 007063	Homo sapiens vascular Rab-GAP/TBC-containing (VRP), mRNA
NM 007027	Homo sapiens topoisomerase (DNA) II binding protein (TOPBP1), mRNA
NM_006938	Homo sapiens small nuclear ribonucleoprotein D1 polypeptide (16kD) (SNRPD1), mRNA
NM_006937	Homo sapiens SMT3 suppressor of mif two 3 homolog 2 (yeast) (SMT3H2), mRNA
NM 007029	Homo sapiens stathmin-like 2 (STMN2), mRNA
NM_007042	Homo sapiens ribonuclease P (14kD) (RPP14), mRNA  Homo sapiens pyrroline-5-carboxylate reductase 1 (PYCR1), nuclear gene
NM_006907	
ND ( 007050	encoding mitochondrial protein, mRNA
NM_007059	Homo sapiens kaptin (actin binding protein) (KPTN), mRNA
NM_007069	Homo sapiens HRAS-like suppressor 3 (HRASLS3), mRNA
NM_006895	Homo sapiens histamine N-methyltransferase (HNMT), mRNA
NM_007071	Homo sapiens HERV-H LTR-associating 3 (HHLA3), mRNA
NM_007067	Homo sapiens histone acetyltransferase (HBOA), mRNA
NM_007006	Homo sapiens cleavage and polyadenylation specific factor 5, 25 kD subunit (CPSF5), mRNA
NM_007053	Homo sapiens natural killer cell receptor, immunoglobulin superfamily member (BY55), mRNA
NM 006754	Homo sapiens synaptophysin-like protein (SYPL), mRNA
NM 006802	Homo sapiens splicing factor 3a, subunit 3, 60kD (SF3A3), mRNA
NM 006842	Homo sapiens splicing factor 3b, subunit 2, 145kD (SF3B2), mRNA
NM 006834	Homo sapiens RAB32, member RAS oncogene family (RAB32), mRNA
11111 000634	1 Homo sapiens (AD32, memoer (AS oneogene family (CAD32), method

	CDD (C) DATA
NM_006875	Homo sapiens pim-2 oncogene (PIM2), mRNA
NM_006810	Homo sapiens for protein disulfide isomerase-related (PDIR), mRNA  Homo sapiens for protein disulfide isomerase-related (PDIR), mRNA
	Homo sapiens HIRA interacting protein 3 (HIRIP3), mRNA  Homo sapiens HIRA interacting protein 3 (Figure 29 (Clorf29) mRNA
NM_006820	Homo sapiens chromosome lopen reading frame 29 (Clorf29), mRNA  Homo sapiens chromosome lopen reading frame 29 (Clorf29), mRNA
NM_006848	Homo sapiens thromesome 1 open the racting protein A (DIPA), mRNA  Homo sapiens hepatitis delta antigen-interacting protein A (DIPA), mRNA
NM_006876	Homo sapiens hepatitis della antigen interactional production of the Homo sapiens UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase
	6 (B3GNT6), mRNA
NM_006653	6 (B3GN16), mRNA  Homo sapiens suc1-associated neurotrophic factor target 2 (FGFR signalling
_	adaptor) (SNT-2), mRNA
NM 006638	Homo sapiens ribonuclease P, 40kD subunit (RPP40), mRNA
NM 004163	TI and DAR27R member RAS oncogene family (RAB27D), find 11
NM 006713	Homo sapiens RAB27B, memori Te le sur la sur la sapiens activated RNA polymerase II transcription cofactor 4 (PC4),
_	mDNA
NM 006601	Homo sapiens unactive progesterone receptor, 23 kD (P23), mRNA
NM 006675	t-t
NM 006501	Homo saniens myelin-associated oligodendrocyte basic protein (WODI), Madis-
NM_006612	The same family member H (KIFIL), IIIINIA
NM_006567	Homo sapiens kinesin faminy member 16 (KM 20), and the sapiens phenylalanine-tRNA synthetase (FARS1), nuclear gene encoding
14141_000507	1 : 1 : 1::-1
NM_006594	Homo sapiens adaptor-related protein complex 4, beta 1 subunit (AP4B1),
MM_000254	7374
NM 006621	Homo saniens S-adenosylhomocysteine hydrolase-like I (AHCYLI), IIIRNA
NM 006472	Home senions thioredoxin interacting protein (IANIF), IIIXIA
NM 006388	Trans agains HTV-1 Tat interactive protein, by KD (HTATH), interior
	Homo sapiens serine/threonine kinase 3 (STE20 homolog, yeast) (STK3),
NM_006281	mDNA
NT 006401	Homo sapiens acidic protein rich in leucines (SSP29), mRNA
NM_006401	to the TI collising factor SI II / (SI II / ), MKNA
NM_006425	Homo sapiens step it splicing factor the (effect), and the Homo sapiens solute carrier family 9 (sodium/hydrogen exchanger), isoform 6
NM_006359	(CI COA6) mRNA
ND4 006228	The DNIA hinding motif protein 14 (RBM14), mKNA
NM_006328	Homo sapiens RNA blidding flottly protein 1 (2007) Homo sapiens polymerase (RNA) III (DNA directed) polypeptide F (39 kD)
NM_006466	mornany Dala
27.5.006467	Homo sapiens polymerase (RNA) III (DNA directed) (32kD) (RPC32), mRNA
NM 006467	Homo sapiens ribonuclease HI, large subunit (RNASEHI), mRNA
NM_006397	Homo sapiens Producted Tri, large substance (RCL), mRNA  Homo sapiens putative c-Myc-responsive (RCL), mRNA
NM_006443	Homo sapiens RAN binding protein 8 (RANBP8), mRNA
NM_006390	Homo sapiens RAIN binding protein 8 (RAINDIO), made
NM_006256	Homo sapiens protein kinase C-like 2 (PRKCL2), mRNA  Homo sapiens protein kinase C-delta (PRKCD), mRNA
NM_006254	Homo sapiens protein kinase C, delta (PRKCD), mRNA  Homo sapiens protein kinase C, delta (PRKCD), mRNA
NM_006229	Homo sapiens pancreatic lipase-related protein 1 (PNLIPRP1), mRNA  Homo sapiens CDP-diacylglycerolinositol 3-phosphatidyltransferase
NM_006319	Homo sapiens CDP-diacylgiycerolmostiol 3-phosphatidyttambretase
	(phosphatidylinositol synthase) (CDIPT), mRNA  (phosphatidylinositol synthase) (CDIPT), mRNA  (phosphatidylinositol synthase) (CDIPT), mRNA
NM_006219	Homo sapiens phosphoinositide-3-kinase, catalytic, beta polypeptide (PIK3CB),
	mRNA
NM 006346	Homo sapiens progesterone-induced blocking factor 1 (PIBF1), mRNA
NM_006473	Homo sapiens TAF6-like RNA polymerase II, p300/CBP-associated factor
_	(DCAT)
NM 006396	Tierro coniona Giogran's syndrome/scleroderma autoantigen i (555CAI), micra
NM 006428	Homo sapiens melanoma-associated antigen recognised by cytotoxic 1
	$1 \cdot 1 \cdot \dots \cdot $
	II was regions actochlast energific factor 2 (fasciclin I-like) (OSF-2), micha
NM 006475	Homo sapiens nucleolar protein 5A (56kD with KKE/D repeat) (NOL5A),

	mRNA
NM 006417	Homo sapiens interferon-induced, hepatitis C-associated microtubular aggregate
_	protein (44kD) (MTAP44), mRNA
NM 006405	Homo sapiens transmembrane 9 superfamily member 1 (TM9SF1), mRNA
NM 006471	Homo sapiens myosin, light polypeptide, regulatory, non-sarcomeric (20kD)
_	(MLCB), mRNA
NM 006152	Homo sapiens lymphoid-restricted membrane protein (LRMP), mRNA
NM 006460	Homo sapiens HMBA-inducible (HIS1), mRNA
NM_006365	Homo sapiens transcriptional activator of the c-fos promoter (CROC4), mRNA
NM 006135	Homo sapiens capping protein (actin filament) muscle Z-line, alpha 1
	(CAPZA1), mRNA
NM_006086	Homo sapiens tubulin, beta, 4 (TUBB4), mRNA
NM_005761	Homo sapiens plexin C1 (PLXNC1), mRNA
NM_005724	Homo sapiens tetraspan 3 (TSPAN-3), mRNA
NM_005646	Homo sapiens TAR (HIV) RNA binding protein 1 (TARBP1), mRNA
NM_005819	Homo sapiens syntaxin 6 (STX6), mRNA
NM_005866	Homo sapiens sigma receptor (SR31747 binding protein 1) (SR-BP1), mRNA
NM_005842	Homo sapiens sprouty homolog 2 (Drosophila) (SPRY2), mRNA
NM_005626	Homo sapiens splicing factor, arginine/serine-rich 4 (SFRS4), mRNA
NM_005770	Homo sapiens small EDRK-rich factor 2 (SERF2), mRNA
NM_005805	Homo sapiens 26S proteasome-associated pad1 homolog (POH1), mRNA
NM_005746	Homo sapiens pre-B-cell colony-enhancing factor (PBEF), mRNA
NM_005869	Homo sapiens serologically defined colon cancer antigen 10 (SDCCAG10),
	mRNA .
NM_005787	Homo sapiens Not56 (D. melanogaster)-like protein (NOT56L), mRNA
NM_005792	Homo sapiens M-phase phosphoprotein 6 (MPHOSPH6), mRNA
NM_005693	Homo sapiens nuclear receptor subfamily 1, group H, member 3 (NR1H3), mRNA
NM_005799	Homo sapiens PDZ domain protein (Drosophila inaD-like) (INADL), mRNA
NM_005713	Homo sapiens collagen, type IV, alpha 3 (Goodpasture antigen) binding protein (COL4A3BP), transcript variant 1, mRNA
NM 005878	Homo sapiens trinucleotide repeat containing 3 (TNRC3), mRNA
NM 005875	Homo sapiens translation factor sui1 homolog (GC20), mRNA
NM_005838	Homo sapiens glycine-N-acyltransferase (GLYAT), nuclear gene encoding mitochondrial protein, mRNA
NM_005754	Homo sapiens Ras-GTPase-activating protein SH3-domain-binding protein (G3BP), mRNA
NM_005764	Homo sapiens epithelial protein up-regulated in carcinoma, membrane associated
	protein 17 (DD96), mRNA
NM_005694	Homo sapiens COX17 homolog, cytochrome c oxidase assembly protein (yeast)
	(COX17), nuclear gene encoding mitochondrial protein, mRNA
NM_005506	Homo sapiens CD36 antigen (collagen type I receptor, thrombospondin
37.5 005001	receptor)-like 2 (lysosomal integral membrane protein II) (CD36L2), mRNA
NM_005881	Homo sapiens branched chain alpha-ketoacid dehydrogenase kinase (BCKDK), mRNA
NM_005718	Homo sapiens actin related protein 2/3 complex, subunit 4 (20 kD) (ARPC4), mRNA
NM_005717	Homo sapiens actin related protein 2/3 complex, subunit 5 (16 kD) (ARPC5), mRNA
NM_005829	Homo sapiens adaptor-related protein complex 3, sigma 2 subunit (AP3S2), mRNA
NM 005814	Homo sapiens glycoprotein A33 (transmembrane) (GPA33), mRNA
1111 003014	Trains supraise Bij coprotein rass (mandinomorano) (or ress); madri

NM_005406	Homo sapiens Rho-associated, coiled-coil containing protein kinase 1 (ROCK1), mRNA
NM_005399	Homo sapiens protein kinase, AMP-activated, beta 2 non-catalytic subunit
	(PRKAB2), mRNA
NM_005396	Homo sapiens pancreatic lipase-related protein 2 (PNLIPRP2), mRNA
NM_005489	Homo sapiens SH2 domain-containing 3C (SH2D3C), mRNA Homo sapiens frequently rearranged in advanced T-cell lymphomas (FRAT1),
NM_005479	mRNA
NM 005154	Homo sapiens ubiquitin specific protease 8 (USP8), mRNA
NM_005066	Homo sapiens splicing factor proline/glutamine rich (polypyrimidine tract binding protein associated) (SFPQ), mRNA
NM_005123	Homo sapiens nuclear receptor subfamily 1, group H, member 4 (NR1H4),
NM 005046	Homo sapiens kallikrein 7 (chymotryptic, stratum corneum) (KLK7), mRNA
NM 005030	Homo sapiens polo-like kinase (Drosophila) (PLK), mRNA
NM 005014	Home conjens esteemodulin (OMD) mRNA
NM_005003	Homo sapiens NADH dehydrogenase (ubiquinone) 1, alpha/beta subcomplex, 1
NM_004941	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 8 (RNA helicase) (DDX8), mRNA
NM 004205	Homo sapiens ubiquitin specific protease 2 (USP2), mRNA
	Homo sapiens prp28, U5 snRNP 100 kd protein (U5-100K), mRNA
NM_004818	Homo sapiens TRF-proximal protein (TRFP), mRNA
NM_004275	Homo sapiens Homer, neuronal immediate early gene, 1B (SYN47), mRNA
NM_004272	Homo conjens syntaxin 3A (STX3A), mRNA
NM_004177 NM_004719	Homo sapiens splicing factor, arginine/serine-rich 2, interacting protein (SFRS2IP), mRNA
NM_004175	Homo sapiens small nuclear ribonucleoprotein D3 polypeptide (18kD)
NM_004592	Homo sapiens splicing factor, arginine/serine-rich 8 (suppressor-of-white-apricot
NM_004799	Homo sapiens MAD, mothers against decapentaplegic homolog (Drosophila) interacting protein, receptor activation anchor (MADHIP), transcript variant 3, mRNA
NM 004875	Homo sapiens RNA polymerase I subunit (RPA40), mRNA
NM 004292	Homo saniens ras inhibitor (RIN1), mRNA
NM 004815	Homo sapiens PTPL1-associated RhoGAP 1 (PARG1), mRNA
NM 004772	Homo saniens P311 protein (P311), mRNA
NM_004553	Homo sapiens NADH dehydrogenase (ubiquinone) Fe-S protein 6 (13kD)
NM_004549	Homo sapiens NADH dehydrogenase (ubiquinone) 1, subcomplex unknown, 2 (14.5kD, B14.5b) (NDUFC2), mRNA
NM_004271	Homo sapiens MD-1 RP105-associated (MD-1), mRNA
NM_004672	Homo sapiens mitogen-activated protein kinase kinase 6 (MAP3K6),
NM_004828	Homo sapiens lymphocyte antigen 95 (activating NK-receptor; NK-p44) (LY95
NM_004735	Homo sapiens leucine rich repeat (in FLII) interacting protein 1 (LRRFIP1), mRNA
NM 004811	Homo sapiens leupaxin (LPXN), mRNA
NM 004511	Homo saniens kinesin family member 5C (KIF5C), mRNA
NM 004905	- landing clutothione nerovidace

	acidic calcium-independent phospholipase A2) (KIAA0106), mRNA
NM 004770	Homo sapiens potassium voltage-gated channel, Shab-related subfamily, member
	2 (KCNB2), mRNA
NM_004848	Homo sapiens basement membrane-induced gene (ICB-1), mRNA
NM_004763	Homo sapiens integrin cytoplasmic domain-associated protein 1 (ICAP-1A), transcript variant 1, mRNA
NM_004814	Homo sapiens U5 snRNP-specific 40 kDa protein (hPrp8-binding) (HPRP8BP), mRNA
NM 004839	Homo sapiens Homer, neuronal immediate early gene, 2 (HOMER-2B), mRNA
NM 004684	Homo sapiens SPARC-like 1 (mast9, hevin) (SPARCL1), mRNA
NM_004832	Homo sapiens glutathione-S-transferase like; glutathione transferase omega (GSTTLp28), mRNA
NM_004486	Homo sapiens golgi autoantigen, golgin subfamily a, 2 (GOLGA2), mRNA
NM 004125	Homo sapiens guanine nucleotide binding protein 10 (GNG10), mRNA
NM_004483	Homo sapiens glycine cleavage system protein H (aminomethyl carrier) (GCSH), mRNA
NM_004767	Homo sapiens endothelin type b receptor-like protein 2 (ET(B)R-LP-2), mRNA
NM_004440	Homo sapiens EphA7 (EPHA7), mRNA
NM_004757	Homo sapiens small inducible cytokine subfamily E, member 1 (endothelial monocyte-activating) (SCYE1), mRNA
NM_004427	Homo sapiens early development regulator 2 (polyhomeotic 2 homolog) (EDR2), mRNA
NM_004422	Homo sapiens dishevelled, dsh homolog 2 (Drosophila) (DVL2), mRNA
NM_004416	Homo sapiens deltex homolog 1 (Drosophila) (DTX1), mRNA
NM_004073	Homo sapiens cytokine-inducible kinase (CNK), mRNA
NM_004365	Homo sapiens centrin, EF-hand protein, 3 (CDC31 homolog, yeast) (CETN3), mRNA
NM_004680	Homo sapiens chromodomain protein, Y chromosome, 1 (CDY1), mRNA
NM_004291	Homo sapiens cocaine- and amphetamine-regulated transcript (CART), mRNA
NM_004330	Homo sapiens BCL2/adenovirus E1B 19kD interacting protein 2 (BNIP2), mRNA
NM_004024	Homo sapiens activating transcription factor 3 (ATF3), mRNA
NM_001177	Homo sapiens ADP-ribosylation factor-like 1 (ARL1), mRNA
NM_001545	Homo sapiens immature colon carcinoma transcript 1 (ICT1), mRNA
NM_001533	Homo sapiens heterogeneous nuclear ribonucleoprotein L (HNRPL), mRNA
NM_001509	Homo sapiens glutathione peroxidase 5 (epididymal androgen-related protein) (GPX5), transcript variant 1, mRNA
NM_001349	Homo sapiens aspartyl-tRNA synthetase (DARS), mRNA
NM_001329	Homo sapiens C-terminal binding protein 2 (CTBP2), transcript variant 1, mRNA
NM_000082	Homo sapiens Cockayne syndrome 1 (classical) (CKN1), mRNA
NM_001277	Homo sapiens choline kinase (CHK), mRNA
NM 001087	Homo sapiens angio-associated, migratory cell protein (AAMP), mRNA
NM 003999	Homo sapiens oncostatin M receptor (OSMR), mRNA
NM_003904	Homo sapiens zinc finger protein 259 (ZNF259), mRNA
NM_003385	Homo sapiens visinin-like 1 (VSNL1), mRNA
NM_003348	Homo sapiens ubiquitin-conjugating enzyme E2N (UBC13 homolog, yeast) (UBE2N), mRNA
NM_003341	Homo sapiens ubiquitin-conjugating enzyme E2E 1 (UBC4/5 homolog, yeast) (UBE2E1), mRNA
NM_003339	Homo sapiens ubiquitin-conjugating enzyme E2D 2 (UBC4/5 homolog, yeast) (UBE2D2), mRNA

apiens UDP-N-acteylglucosamine pyrophosphorylase 1 (UAP1), mRNA apiens transient receptor potential cation channel, subfamily C, member C3), mRNA apiens tyrosylprotein sulfotransferase 1 (TPST1), mRNA apiens tankyrase, TRF1-interacting ankyrin-related ADP-ribose ase (TNKS), mRNA apiens syntaxin 7 (STX7), mRNA apiens syntaxin 5A (STX5A), mRNA apiens syntaxin 11 (STX11), mRNA apiens signal recognition particle 9kD (SRP9), mRNA apiens signal recognition particle 54kD (SRP54), mRNA apiens serum response factor (c-fos serum response element-binding ption factor) (SRF), mRNA apiens sorting nexin 3 (SNX3), mRNA sapiens small nuclear ribonucleoprotein polypeptide G (SNRPG), mRNA
apiens tyrosylprotein sulfotransferase 1 (TPST1), mRNA apiens tyrosylprotein sulfotransferase 1 (TPST1), mRNA apiens tankyrase, TRF1-interacting ankyrin-related ADP-ribose ase (TNKS), mRNA apiens syntaxin 7 (STX7), mRNA apiens syntaxin 5A (STX5A), mRNA apiens syntaxin 11 (STX11), mRNA apiens signal recognition particle 9kD (SRP9), mRNA apiens signal recognition particle 54kD (SRP54), mRNA apiens serum response factor (c-fos serum response element-binding ption factor) (SRF), mRNA apiens sorting nexin 3 (SNX3), mRNA
apiens tankyrase, TRF1-interacting ankyrin-related ADF-Hoose ase (TNKS), mRNA apiens syntaxin 7 (STX7), mRNA apiens syntaxin 5A (STX5A), mRNA apiens syntaxin 11 (STX11), mRNA apiens signal recognition particle 9kD (SRP9), mRNA apiens signal recognition particle 54kD (SRP54), mRNA apiens serum response factor (c-fos serum response element-binding ption factor) (SRF), mRNA apiens sorting nexin 3 (SNX3), mRNA
ase (TNKS), mRNA apiens syntaxin 7 (STX7), mRNA apiens syntaxin 5A (STX5A), mRNA apiens syntaxin 11 (STX11), mRNA apiens signal recognition particle 9kD (SRP9), mRNA apiens signal recognition particle 54kD (SRP54), mRNA apiens serum response factor (c-fos serum response element-binding ption factor) (SRF), mRNA apiens sorting nexin 3 (SNX3), mRNA
ase (TNKS), mRNA apiens syntaxin 7 (STX7), mRNA apiens syntaxin 5A (STX5A), mRNA apiens syntaxin 11 (STX11), mRNA apiens signal recognition particle 9kD (SRP9), mRNA apiens signal recognition particle 54kD (SRP54), mRNA apiens serum response factor (c-fos serum response element-binding ption factor) (SRF), mRNA apiens sorting nexin 3 (SNX3), mRNA
apiens syntaxin 5A (STX5A), mRNA apiens syntaxin 11 (STX11), mRNA apiens signal recognition particle 9kD (SRP9), mRNA apiens signal recognition particle 54kD (SRP54), mRNA apiens serum response factor (c-fos serum response element-binding ption factor) (SRF), mRNA apiens sorting nexin 3 (SNX3), mRNA
apiens syntaxin 11 (STX11), mRNA apiens signal recognition particle 9kD (SRP9), mRNA apiens signal recognition particle 54kD (SRP54), mRNA apiens serum response factor (c-fos serum response element-binding ption factor) (SRF), mRNA sapiens sorting nexin 3 (SNX3), mRNA
apiens signal recognition particle 9kD (SRP9), mRNA apiens signal recognition particle 54kD (SRP54), mRNA apiens serum response factor (c-fos serum response element-binding ption factor) (SRF), mRNA apiens sorting nexin 3 (SNX3), mRNA
apiens signal recognition particle 54kD (SRP34), mkNA apiens serum response factor (c-fos serum response element-binding ption factor) (SRF), mRNA sapiens sorting nexin 3 (SNX3), mRNA sapiens small nuclear ribonucleoprotein polypeptide G (SNRPG), mRNA
apiens serum response factor (c-tos serum response element-oniding ption factor) (SRF), mRNA sapiens sorting nexin 3 (SNX3), mRNA
ption factor) (SRF), mRNA sapiens sorting nexin 3 (SNX3), mRNA sapiens small nuclear ribonucleoprotein polypeptide G (SNRPG), mRNA
sapiens sorting nexin 3 (SNX3), mRNA
sapiens sorting nexin 3 (SNX3), mRNA
amions small nuclear ribonucleoprotein polypeptide G (SING G), IIIGAR
apiens small (CAMDIC) mPNA
sapiens small nuclear ribonucleoprotein polypeptide C (SNRPC), mRNA
sapiens sphingomyelin phosphodiesterase 2, neutral memorane (neutral
lings) (SMDD2) mRNA
sapiens solute carrier family 22 (organic cation transporter), member 4
2 \ \ \ \ mPN \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \
sapiens sialyltransferase 4A (beta-galactosidase alpha-2,3-
foreso (SIATAA) mRNA
sapiens ribosomal protein S6 kinase, 70kD, polypeptide 2 (RPS6KB2),
sapiens RTC domain containing 1 (RTCD1), mRNA
conjons ribonuclease RNase A family, 4 (RNASE4), mrina
sapiens receptor (TNFRSF)-interacting serine-threonine kinase 1
1\DNA
sapiens RNA binding motif, single stranded interacting protein 2
C2) mDNA
sapiens RAP2B, member of RAS oncogene family (RAP2B), mRNA
and any olin protein zero-like I (MPZLII), mKNA
sapiens invertif protein zero inte 1 (1222), sapiens proteasome (prosome, macropain) 26S subunit, non-ATPase, 3
D3) mRNA
senions protease, serine 3 (trypsin 3) (PRSS3), mRNA
iona mitogen activated protein kinase kinase o (MACAN), illicita
copiens mitogen-activated protein kinase 13 (MAPK 13), mkina
sapiens mitogen activated protein kinase-activated protein kinase 5
DV ADVS) mDNA
soniens protein phosphatase 2 (formerly 2A), regulatory subunit B (FK
Inha isoform and (PR 130), beta isoform (PPP2R3), mRNA
o sapiens PTPRF interacting protein, binding protein 1 (liprin beta 1)
IPPI) mPNIA
saniens protein tyrosine phosphatase, receptor type, i polypeptide
DE) interacting protein (lingin), alpha I (PPFIAI), MKINA
a comions not merase (DNA-directed), alpha (70KD) (POLAZ), IIIXVA
o sapiens polymyositis/scleroderma autoantigen 2 (100kD) (PMSCL2),
A
o sapiens putative receptor protein (PMI), mRNA
o sapiens plastin 1 (I isoform) (PLS1), mRNA
o sapiens plastin 1 (I isoform) (PLS1), mRNA o sapiens pleckstrin (PLEK), mRNA o sapiens phosphatidylinositol-4-phosphate 5-kinase, type II, beta

	(mmsyap)DNA
37.6.002.620	(PIP5K2B), mRNA
NM_003629	Homo sapiens phosphoinositide-3-kinase, regulatory subunit, polypeptide 3 (p55,
37.5.002.640	gamma) (PIK3R3), mRNA
NM_002649	Homo sapiens phosphoinositide-3-kinase, catalytic, gamma polypeptide
	(PIK3CG), mRNA
NM_002624	Homo sapiens prefoldin 5 (PFDN5), mRNA
NM_003846	Homo sapiens peroxisomal biogenesis factor 11B (PEX11B), mRNA
NM_002617	Homo sapiens peroxisome biogenesis factor 10 (PEX10), mRNA
NM_002611	Homo sapiens pyruvate dehydrogenase kinase, isoenzyme 2 (PDK2), mRNA
NM_000923	Homo sapiens phosphodiesterase 4C, cAMP-specific (phosphodiesterase E1 dunce homolog, Drosophila) (PDE4C), mRNA
NM 002599	Homo sapiens phosphodiesterase 2A, cGMP-stimulated (PDE2A), mRNA
NM 002504	Homo sapiens nuclear transcription factor, X-box binding 1 (NFX1), mRNA
NM_002482	Homo sapiens nuclear autoantigenic sperm protein (histone-binding) (NASP), mRNA
NM_003826	Homo sapiens N-ethylmaleimide-sensitive factor attachment protein, gamma (NAPG), mRNA
NM 002465	Homo sapiens myosin binding protein C, slow type (MYBPC1), mRNA
NM 002461	Homo sapiens mevalonate (diphospho) decarboxylase (MVD), mRNA
NM 002461 NM 003676	Homo sapiens degenerative spermatocyte homolog, lipid desaturase (Drosophila)
	(DEGS), mRNA
NM_002307	Homo sapiens lectin, galactoside-binding, soluble, 7 (galectin 7) (LGALS7), mRNA
NM 002271	Homo sapiens karyopherin (importin) beta 3 (KPNB3), mRNA
NM 002270	Homo sapiens karyopherin (importin) beta 2 (KPNB2), mRNA
NM 002214	Homo sapiens integrin, beta 8 (ITGB8), mRNA
NM_002204	Homo sapiens integrin, alpha 3 (antigen CD49C, alpha 3 subunit of VLA-3 receptor) (ITGA3), transcript variant a, mRNA
NM 001560	Homo sapiens interleukin 13 receptor, alpha 1 (IL13RA1), mRNA
NM 002163	Homo sapiens interferon consensus sequence binding protein 1 (ICSBP1),
	mRNA
NM_002156	Homo sapiens heat shock 60kD protein 1 (chaperonin) (HSPD1), mRNA
NM_002149	Homo sapiens hippocalcin-like 1 (HPCAL1), mRNA
NM_003947	Homo sapiens huntingtin-associated protein interacting protein (duo) (HAPIP), mRNA
NM_003665	Homo sapiens ficolin (collagen/fibrinogen domain containing) 3 (Hakata antigen) (FCN3), mRNA
NM 000842	Homo sapiens glutamate receptor, metabotropic 5 (GRM5), mRNA
NM_002053	Homo sapiens guanylate binding protein 1, interferon-inducible, 67kD (GBP1), mRNA
NM 001482	Homo sapiens glycine amidinotransferase (L-arginine:glycine
14141_001402	amidinotransferase) (GATM), mRNA
NM 002044	Homo sapiens galactokinase 2 (GALK2), mRNA
NM 001417	Homo sapiens eukaryotic translation initiation factor 4B (EIF4B), mRNA
NM 003758	Homo sapiens eukaryotic translation initiation factor 3, subunit 1 (alpha, 35kD)
	(EIF3S1), mRNA
NM_001404	Homo sapiens eukaryotic translation elongation factor 1 gamma (EEF1G), mRNA
NM_001960	Homo sapiens eukaryotic translation elongation factor 1 delta (guanine nucleotide exchange protein) (EEF1D), mRNA
NM_003792	Homo sapiens endothelial differentiation-related factor 1 (EDF1), mRNA
NM 003974	Homo sapiens docking protein 2, 56kD (DOK2), mRNA
141VI_0039/4	I from sapiens docking protein 2, 30kD (DOK2), niktva

	The state of the s
NM_003586	Homo sapiens double C2-like domains, alpha (DOC2A), mRNA
	Homo sapiens double C2-like dollaris, apple (2003), Homo sapiens corticotropin releasing hormone receptor 2 (CRHR2), mRNA
	Homo sapiens carboxypeptidase E (CPE), mRNA
NM 001782	Homo sapiens CD72 antigen (CD72), mRNA  TCP1 submit 6A (zeta 1) (CCT6A)
	Homo sapiens CD72 anugen (CD72), indexer  Homo sapiens chaperonin containing TCP1, subunit 6A (zeta 1) (CCT6A),
	mRNA Homo sapiens Ca2+-dependent activator protein for secretion (CADPS), mRNA
NM 003986	Homo saniens butvrobetaine (gamma), 2-oxogiutarate dioxygenase (gamma
IAIM_002300	but we betaine hydroxylase)   (BBOXI), MKINA
NM 001674	XI conjugation transcription factor 3 (A1F3), mkNA
NM 001074 NM 001173	Homo saniens Rho GTPase activating protein 5 (ARHOAI 5), micro
NM 025065	Homo saniens RNA processing factor 1 (RPF1), mRNA
NM 024907	They protein FRG4 (FR(i4), MKNA
	Trienhosphate 3-kinase C (III RC), michair
NM_025194 NM_014203	Homo sapiens adaptor-related protein complex 2, alpha 1 subunit (AP2A1),
14141_01 1205	mRNA
NM 130786	11 1 D. Languagetoin (A1RG) mRNA
NM 031482	Homo saniens hypothetical protein DKFZp58610418 (DKFZF58010418), interior
NM 015419	TI-ma conjone adlican (DKF7n56411922), mKNA
NM 015683	Try ham athetical protein ((1,0)NE/4943), IIINNA
NM 015638	TI chromosome 20 onen reading frame 100 (C20011100), mid di
NM 080737	Home conjens symantota omin-like 4 (granuphilin-a) (STIL4), mixtx
NM 080723	Homo saniens vesicular membrane protein p24 (VMP), mknA
NM_080723	TT : NEDDS conjugating enzyme (NCE2), MKINA
	1
NM_080668	Homo sapiens similar to RIKEN cDNA 2600001A11 gene (LOC112840),
NM_080666	
NM 080663	$+$ $\pm$
NM 080661	The transfer of the control of the c
NM 080658	T == 0 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
NM_080656	Homo sapiens similar to RIKEN cDNA A430101B06 gene (MGC13017),
MM_080030	1
ND 6 000651	initianto DIVEN cDNA 1810038N03 gene (MGC9890), IIINA
NM_080651	Thems comions similar to RIKEN CONA 3/30421E16 gene (MGC14790); 1122
NM_080650	Tax in the function protein 4 (peripheral) (1)P4), IIINIA
NM_080604	Homo sapiens tight junction protein 4 (po acid transporter (VIAAT), mRNA  Homo sapiens vesicular inhibitory amino acid transporter (VIAAT), mRNA
NM_080552	Homo sapiens aquaporin 10 (AQP10), mRNA  Homo sapiens aquaporin 10 (AQP10), mRNA
NM_080429	Homo sapiens axonemal dynein heavy chain 7 (DNAH7), mRNA  Homo sapiens axonemal dynein heavy chain 7 (DNAH7), mRNA
NM_018897	Homo sapiens autism-related protein 1 (KIAA0442), mRNA
NM_015570	
NM_015132	Homo sapiens sorting flexin 13 (SNX13), find 17  Homo sapiens similar to constitutive photomorphogenic protein 1 (Arabidopsis)
NM_022457	1
NTM 020650	(FLJ10416), mRNA  Homo sapiens putative ankyrin-repeat containing protein (DKFZP564D166),
NM_030658	mRNA
NM 058229	Harris ganions E hav only protein 32 (FBXO32), mRNA
NM 058188	Transport of the strong spreading frame 6/ (C210107), HIGNA
NM 058187	Home conjens chromosome 21 open reading frame 03 (C2101103), find 121
I TATAT ODGIGA	Homo saniens ING1-like tumor suppressor protein (ING1-like), mid 17
NM_058171	Homo sablens ubiquitin confugating charing charing
NM_058171 NM_058167	Homo saniens centaurin delta 2 (CENTD2), mRINA
NM_058171	Homo sapiens centaurin, delta 2 (CENTD2), mRNA

NM_054108	Homo sapiens H-rev107-like protein 5 (HRLP5), mRNA
NM_020794	Homo sapiens densin-180 (KIAA1365), mRNA
NM_054032	Homo sapiens G protein-coupled receptor MRGX4 (MRGX4), mRNA
NM 054031	Homo sapiens G protein-coupled receptor MRGX3 (MRGX3), mRNA
NM 054030	Homo sapiens G protein-coupled receptor MRGX2 (MRGX2), mRNA
NM 054023	Homo sapiens uteroglobin-related protein 1 (UGRP1), mRNA
NM 054024	Homo sapiens melanoma inhibitory activity protein 2 (MIA2), mRNA
NM 031946	Homo sapiens centaurin, gamma 3 (CENTG3), mRNA
NM 052860	Homo sapiens kruppel-like zinc finger protein (ZNF300), mRNA
NM 053054	Homo sapiens cation channel of sperm (CATSPER), mRNA
NM 053053	Homo sapiens SPT3-associated factor 42 (STAF42), mRNA
NM 053048	Homo sapiens hypothetical protein MGC16384 (MGC16384), mRNA
NM 053047	Homo sapiens hypothetical protein MGC16063 (MGC16063), mRNA
NM 053040	Homo sapiens PNAS-123 (LOC85028), mRNA
NM 053039	Homo sapiens UDP glycosyltransferase 2 family, polypeptide B28 (UGT2B28),
	mRNA
NM 053001	Homo sapiens odd-skipped-related 2A protein (OSR2), mRNA
NM 052997	Homo sapiens breast cancer antigen NY-BR-1 (NY-BR-1), mRNA
NM 052971	Homo sapiens liver-expressed antimicrobial peptide 2 (LEAP-2), mRNA
NM 052956	Homo sapiens medium-chain acyl-CoA synthetase (MACS1), mRNA
NM 052942	Homo sapiens guanylate binding protein 5 (GBP5), mRNA
NM 052931	Homo sapiens activating NK receptor (KALI), mRNA
NM 052879	Homo sapiens c-Mpl binding protein (LOC113251), mRNA
NM 030928	Homo sapiens DNA replication factor (CDT1), mRNA
NM 025185	Homo sapiens putative ankyrin-repeat containing protein (DKFZP564D166),
10025105	mRNA
NM 015179	Homo sapiens KIAA0690 protein (KIAA0690), mRNA
NM 033626	Homo sapiens JM11 protein (JM11), mRNA
NM 022735	Homo sapiens golgi phosphoprotein 1 (GOLPH1), mRNA
NM 033547	Homo sapiens hypothetical gene MGC16733 similar to CG12113 (MGC16733),
_	mRNA
NM_032268	Homo sapiens nerve injury gene 283 (NIN283), mRNA
NM_016167	
	Homo sapiens retinoic acid repressible protein (RARG-1), mRNA
NM_033414	Homo sapiens retinoic acid repressible protein (RARG-1), mRNA  Homo sapiens hypothetical protein MGC17552 (MGC17552), mRNA
NM_033414 NM_016336	Homo sapiens hypothetical protein MGC17552 (MGC17552), mRNA
	Homo sapiens hypothetical protein MGC17552 (MGC17552), mRNA Homo sapiens non-canonical ubquitin conjugating enzyme 1 (NCUBE1), mRNA
NM_016336	Homo sapiens hypothetical protein MGC17552 (MGC17552), mRNA Homo sapiens non-canonical ubquitin conjugating enzyme 1 (NCUBE1), mRNA Homo sapiens hypothetical gene ZD52F10 (ZD52F10), mRNA
NM_016336 NM_033317	Homo sapiens hypothetical protein MGC17552 (MGC17552), mRNA Homo sapiens non-canonical ubquitin conjugating enzyme 1 (NCUBE1), mRNA
NM 016336 NM 033317 NM 033266	Homo sapiens hypothetical protein MGC17552 (MGC17552), mRNA Homo sapiens non-canonical ubquitin conjugating enzyme 1 (NCUBE1), mRNA Homo sapiens hypothetical gene ZD52F10 (ZD52F10), mRNA Homo sapiens ER to nucleus signalling 2 (ERN2), mRNA
NM 016336 NM 033317 NM 033266 NM 031955 NM 033210	Homo sapiens hypothetical protein MGC17552 (MGC17552), mRNA  Homo sapiens non-canonical ubquitin conjugating enzyme 1 (NCUBE1), mRNA  Homo sapiens hypothetical gene ZD52F10 (ZD52F10), mRNA  Homo sapiens ER to nucleus signalling 2 (ERN2), mRNA  Homo sapiens NYD-SP12 protein (NYD-SP12), mRNA  Homo sapiens hypothetical protein FLJ14855 (FLJ14855), mRNA
NM 016336 NM 033317 NM 033266 NM 031955	Homo sapiens hypothetical protein MGC17552 (MGC17552), mRNA Homo sapiens non-canonical ubquitin conjugating enzyme 1 (NCUBE1), mRNA Homo sapiens hypothetical gene ZD52F10 (ZD52F10), mRNA Homo sapiens ER to nucleus signalling 2 (ERN2), mRNA Homo sapiens NYD-SP12 protein (NYD-SP12), mRNA
NM 016336 NM 033317 NM 033266 NM 031955 NM 033210 NM 033211	Homo sapiens hypothetical protein MGC17552 (MGC17552), mRNA Homo sapiens non-canonical ubquitin conjugating enzyme 1 (NCUBE1), mRNA Homo sapiens hypothetical gene ZD52F10 (ZD52F10), mRNA Homo sapiens ER to nucleus signalling 2 (ERN2), mRNA Homo sapiens NYD-SP12 protein (NYD-SP12), mRNA Homo sapiens hypothetical protein FLJ14855 (FLJ14855), mRNA Homo sapiens hypothetical gene supported by AF038182; BC009203 (LOC90355), mRNA
NM 016336 NM 033317 NM 033266 NM 031955 NM 033210 NM 033211	Homo sapiens hypothetical protein MGC17552 (MGC17552), mRNA Homo sapiens non-canonical ubquitin conjugating enzyme 1 (NCUBE1), mRNA Homo sapiens hypothetical gene ZD52F10 (ZD52F10), mRNA Homo sapiens ER to nucleus signalling 2 (ERN2), mRNA Homo sapiens NYD-SP12 protein (NYD-SP12), mRNA Homo sapiens hypothetical protein FLJ14855 (FLJ14855), mRNA Homo sapiens hypothetical gene supported by AF038182; BC009203 (LOC90355), mRNA Homo sapiens small heat shock protein B9 (HspB9), mRNA
NM 016336 NM 033317 NM 033266 NM 031955 NM 033210 NM 033211 NM 033194 NM 032122	Homo sapiens hypothetical protein MGC17552 (MGC17552), mRNA Homo sapiens non-canonical ubquitin conjugating enzyme 1 (NCUBE1), mRNA Homo sapiens hypothetical gene ZD52F10 (ZD52F10), mRNA Homo sapiens ER to nucleus signalling 2 (ERN2), mRNA Homo sapiens NYD-SP12 protein (NYD-SP12), mRNA Homo sapiens hypothetical protein FLJ14855 (FLJ14855), mRNA Homo sapiens hypothetical gene supported by AF038182; BC009203 (LOC90355), mRNA Homo sapiens small heat shock protein B9 (HspB9), mRNA Homo sapiens dystrobrevin binding protein 1 (DTNBP1), mRNA
NM 016336 NM 033317 NM 033266 NM 031955 NM 033210 NM 033211 NM 033194 NM 032122 NM 020405	Homo sapiens hypothetical protein MGC17552 (MGC17552), mRNA Homo sapiens non-canonical ubquitin conjugating enzyme 1 (NCUBE1), mRNA Homo sapiens hypothetical gene ZD52F10 (ZD52F10), mRNA Homo sapiens ER to nucleus signalling 2 (ERN2), mRNA Homo sapiens NYD-SP12 protein (NYD-SP12), mRNA Homo sapiens hypothetical protein FLJ14855 (FLJ14855), mRNA Homo sapiens hypothetical gene supported by AF038182; BC009203 (LOC90355), mRNA Homo sapiens small heat shock protein B9 (HspB9), mRNA Homo sapiens dystrobrevin binding protein 1 (DTNBP1), mRNA Homo sapiens tumor endothelial marker 7 precursor (TEM7), mRNA
NM 016336 NM 033317 NM 033266 NM 031955 NM 033210 NM 033211 NM 033194 NM 032122 NM 020405 NM 033115	Homo sapiens hypothetical protein MGC17552 (MGC17552), mRNA Homo sapiens non-canonical ubquitin conjugating enzyme 1 (NCUBE1), mRNA Homo sapiens hypothetical gene ZD52F10 (ZD52F10), mRNA Homo sapiens ER to nucleus signalling 2 (ERN2), mRNA Homo sapiens NYD-SP12 protein (NYD-SP12), mRNA Homo sapiens hypothetical protein FLJ14855 (FLJ14855), mRNA Homo sapiens hypothetical gene supported by AF038182; BC009203 (LOC90355), mRNA Homo sapiens small heat shock protein B9 (HspB9), mRNA Homo sapiens dystrobrevin binding protein 1 (DTNBP1), mRNA Homo sapiens tumor endothelial marker 7 precursor (TEM7), mRNA Homo sapiens hypothetical protein MGC16169 (MGC16169), mRNA
NM 016336 NM 033317 NM 033266 NM 031955 NM 033210 NM 033211 NM 033194 NM 032122 NM 020405 NM 033115 NM 033117	Homo sapiens hypothetical protein MGC17552 (MGC17552), mRNA Homo sapiens non-canonical ubquitin conjugating enzyme 1 (NCUBE1), mRNA Homo sapiens hypothetical gene ZD52F10 (ZD52F10), mRNA Homo sapiens ER to nucleus signalling 2 (ERN2), mRNA Homo sapiens NYD-SP12 protein (NYD-SP12), mRNA Homo sapiens hypothetical protein FLJ14855 (FLJ14855), mRNA Homo sapiens hypothetical gene supported by AF038182; BC009203 (LOC90355), mRNA Homo sapiens small heat shock protein B9 (HspB9), mRNA Homo sapiens dystrobrevin binding protein 1 (DTNBP1), mRNA Homo sapiens tumor endothelial marker 7 precursor (TEM7), mRNA Homo sapiens hypothetical protein MGC16169 (MGC16169), mRNA Homo sapiens hypothetical protein MGC2734 (MGC2734), mRNA
NM 016336 NM 033317 NM 033266 NM 031955 NM 033210 NM 033211 NM 033194 NM 032122 NM 020405 NM 033115 NM 033117 NM 033103	Homo sapiens hypothetical protein MGC17552 (MGC17552), mRNA Homo sapiens non-canonical ubquitin conjugating enzyme 1 (NCUBE1), mRNA Homo sapiens hypothetical gene ZD52F10 (ZD52F10), mRNA Homo sapiens ER to nucleus signalling 2 (ERN2), mRNA Homo sapiens NYD-SP12 protein (NYD-SP12), mRNA Homo sapiens hypothetical protein FLJ14855 (FLJ14855), mRNA Homo sapiens hypothetical gene supported by AF038182; BC009203 (LOC90355), mRNA Homo sapiens small heat shock protein B9 (HspB9), mRNA Homo sapiens dystrobrevin binding protein 1 (DTNBP1), mRNA Homo sapiens tumor endothelial marker 7 precursor (TEM7), mRNA Homo sapiens hypothetical protein MGC16169 (MGC16169), mRNA Homo sapiens hypothetical protein MGC2734 (MGC2734), mRNA Homo sapiens rhophilin-like protein (LOC85415), mRNA
NM 016336 NM 033317 NM 033266 NM 031955 NM 033210 NM 033211 NM 033194 NM 032122 NM 020405 NM 033115 NM 033117 NM 033103 NM 033035	Homo sapiens hypothetical protein MGC17552 (MGC17552), mRNA Homo sapiens non-canonical ubquitin conjugating enzyme 1 (NCUBE1), mRNA Homo sapiens hypothetical gene ZD52F10 (ZD52F10), mRNA Homo sapiens ER to nucleus signalling 2 (ERN2), mRNA Homo sapiens NYD-SP12 protein (NYD-SP12), mRNA Homo sapiens hypothetical protein FLJ14855 (FLJ14855), mRNA Homo sapiens hypothetical gene supported by AF038182; BC009203 (LOC90355), mRNA Homo sapiens small heat shock protein B9 (HspB9), mRNA Homo sapiens dystrobrevin binding protein 1 (DTNBP1), mRNA Homo sapiens tumor endothelial marker 7 precursor (TEM7), mRNA Homo sapiens hypothetical protein MGC16169 (MGC16169), mRNA Homo sapiens hypothetical protein MGC2734 (MGC2734), mRNA Homo sapiens rhophilin-like protein (LOC85415), mRNA Homo sapiens thymic stromal lymphopoietin (TSLP), mRNA
NM 016336 NM 033317 NM 033266 NM 031955 NM 033210 NM 033211 NM 033194 NM 032122 NM 020405 NM 033115 NM 033117 NM 033103	Homo sapiens hypothetical protein MGC17552 (MGC17552), mRNA Homo sapiens non-canonical ubquitin conjugating enzyme 1 (NCUBE1), mRNA Homo sapiens hypothetical gene ZD52F10 (ZD52F10), mRNA Homo sapiens ER to nucleus signalling 2 (ERN2), mRNA Homo sapiens NYD-SP12 protein (NYD-SP12), mRNA Homo sapiens hypothetical protein FLJ14855 (FLJ14855), mRNA Homo sapiens hypothetical gene supported by AF038182; BC009203 (LOC90355), mRNA Homo sapiens small heat shock protein B9 (HspB9), mRNA Homo sapiens dystrobrevin binding protein 1 (DTNBP1), mRNA Homo sapiens tumor endothelial marker 7 precursor (TEM7), mRNA Homo sapiens hypothetical protein MGC16169 (MGC16169), mRNA Homo sapiens hypothetical protein MGC2734 (MGC2734), mRNA Homo sapiens rhophilin-like protein (LOC85415), mRNA Homo sapiens golgi associated, gamma adaptin ear containing, ARF binding
NM 016336 NM 033317 NM 033266 NM 031955 NM 033210 NM 033211 NM 033194 NM 032122 NM 020405 NM 033115 NM 033117 NM 033103 NM 033035 NM 014001	Homo sapiens hypothetical protein MGC17552 (MGC17552), mRNA Homo sapiens non-canonical ubquitin conjugating enzyme 1 (NCUBE1), mRNA Homo sapiens hypothetical gene ZD52F10 (ZD52F10), mRNA Homo sapiens ER to nucleus signalling 2 (ERN2), mRNA Homo sapiens NYD-SP12 protein (NYD-SP12), mRNA Homo sapiens hypothetical protein FLJ14855 (FLJ14855), mRNA Homo sapiens hypothetical gene supported by AF038182; BC009203 (LOC90355), mRNA Homo sapiens small heat shock protein B9 (HspB9), mRNA Homo sapiens dystrobrevin binding protein 1 (DTNBP1), mRNA Homo sapiens tumor endothelial marker 7 precursor (TEM7), mRNA Homo sapiens hypothetical protein MGC16169 (MGC16169), mRNA Homo sapiens hypothetical protein MGC2734 (MGC2734), mRNA Homo sapiens rhophilin-like protein (LOC85415), mRNA Homo sapiens golgi associated, gamma adaptin ear containing, ARF binding protein 3 (GGA3), mRNA
NM 016336 NM 033317 NM 033266 NM 031955 NM 033210 NM 033211 NM 033194 NM 032122 NM 020405 NM 033115 NM 033117 NM 033103 NM 033035	Homo sapiens hypothetical protein MGC17552 (MGC17552), mRNA Homo sapiens non-canonical ubquitin conjugating enzyme 1 (NCUBE1), mRNA Homo sapiens hypothetical gene ZD52F10 (ZD52F10), mRNA Homo sapiens ER to nucleus signalling 2 (ERN2), mRNA Homo sapiens NYD-SP12 protein (NYD-SP12), mRNA Homo sapiens hypothetical protein FLJ14855 (FLJ14855), mRNA Homo sapiens hypothetical gene supported by AF038182; BC009203 (LOC90355), mRNA Homo sapiens small heat shock protein B9 (HspB9), mRNA Homo sapiens dystrobrevin binding protein 1 (DTNBP1), mRNA Homo sapiens tumor endothelial marker 7 precursor (TEM7), mRNA Homo sapiens hypothetical protein MGC16169 (MGC16169), mRNA Homo sapiens hypothetical protein MGC2734 (MGC2734), mRNA Homo sapiens rhophilin-like protein (LOC85415), mRNA Homo sapiens golgi associated, gamma adaptin ear containing, ARF binding

NM_032932	Homo sapiens hypothetical protein MGC11316 (MGC11316), mRNA
NM_032930	Homo sapiens hypothetical protein MGC13040 (MGC13040), mRNA
NM 032918	Homo sapiens RAS-like, estrogen-regulated, growth-inhibitor (RERG), mRNA
NM 032916	Homo sapiens hypothetical protein MGC16279 (MGC16279), mRNA
NM_032907	Homo sapiens hypothetical protein MGC14421 (MGC14421), mRNA
NM_032904	Homo sapiens hypothetical protein MGC14433 (MGC14433), mRNA
NM_032900	Homo sapiens hypothetical protein MGC14258 (MGC14258), mRNA
NM_032895	Homo sapiens hypothetical protein MGC14376 (MGC14376), mRNA
NM_032888	Homo sapiens KIAA1870 protein (KIAA1870), mRNA
NM_032886	Homo sapiens hypothetical protein MGC15912 (MGC15912), mRNA
NM 032884	Homo sapiens hypothetical protein MGC15882 (MGC15882), mRNA
NM_032876	Homo sapiens hypothetical protein MGC15563 (MGC15563), mRNA
NM 032875	Homo sapiens hypothetical protein MGC15482 (MGC15482), mRNA
NM 032874	Homo sapiens hypothetical protein MGC15438 (MGC15438), mRNA
NM_032872	Homo sapiens NADPH oxidase-related, C2 domain-containing protein (JFC1), mRNA
NM_032871	Homo sapiens tumor necrosis factor receptor superfamily, member 19-like
ND4 022966	(TNFRSF19L), mRNA  Hermo coming hymothetical protein EU 114057 (EU 114057) mRNA
NM_032866	Homo sapiens hypothetical protein FLJ14957 (FLJ14957), mRNA
NM_032860	Homo sapiens hypothetical protein FLJ14909 (FLJ14909), mRNA Homo sapiens hypothetical protein FLJ14904 (FLJ14904), mRNA
NM_032858	Homo sapiens AUT-like 1, cysteine endopeptidase (S. cerevisiae) (AUTL1),
NM_032852	mRNA
NM 032848	Homo sapiens hypothetical protein FLJ14827 (FLJ14827), mRNA
NM 032845	Homo sapiens hypothetical protein FLJ14816 (FLJ14816), mRNA
NM 032835	Homo sapiens hypothetical protein FLJ14761 (FLJ14761), mRNA
NM 032824	Homo sapiens hypothetical protein FLJ14/61 (FLJ14/61), mRNA
NM 032823	Homo sapiens hypothetical protein FLJ14675 (FLJ14675), mRNA
NM_032822	Homo sapiens hypothetical protein FLJ14668 (FLJ14668), mRNA
NM 032828	Homo sapiens hypothetical protein FLJ14642 (FLJ14642), mRNA
NM 032804	Homo sapiens hypothetical protein FLJ14547 (FLJ14547), mRNA
NM 032795	Homo sapiens hypothetical protein FLJ14494 (FLJ14494), mRNA
NM 032783	Homo sapiens hypothetical protein FLJ14431 (FLJ14431), mRNA
NM 032766	Homo sapiens hypothetical protein MGC16179 (MGC16179), mRNA
NM 032763	Homo sapiens hypothetical protein MGC16142 (MGC16142), mRNA
NM 032756	Homo sapiens hypothetical protein MGC15668 (MGC15668), mRNA
NM 032744	Homo sapiens hypothetical protein MGC13008 (MGC13008), inRNA  Homo sapiens hypothetical protein MGC12335 (MGC12335), mRNA
NM 032738	Homo sapiens hypothetical protein MGC12333 (MGC12333), inRNA  Homo sapiens hypothetical protein MGC4595 (MGC4595), mRNA
NM 032723	Homo sapiens hypothetical protein MGC4393 (MGC4393), mRNA  Homo sapiens hypothetical protein MGC12760 (MGC12760), mRNA
NM 032720	Homo sapiens hypothetical protein MGC12760 (MGC12760), inRNA  Homo sapiens hypothetical protein MGC10724 (MGC10724), mRNA
NM 032720	Homo sapiens hypothetical protein MGC4643 (MGC4643), mRNA
NM_032712	Homo sapiens hypothetical protein MGC4043 (MGC4043), mRNA  Homo sapiens hypothetical protein MGC13170 (MGC13170), mRNA
NM 032712 NM 032711	Homo sapiens hypothetical protein MGC13170 (MGC13170), mRNA  Homo sapiens hypothetical protein MGC13090 (MGC13090), mRNA
NM 032711	Homo sapiens hypothetical protein MGC13090 (MGC13090), mRNA  Homo sapiens hypothetical protein MGC12966 (MGC12966), mRNA
NM 032705	Homo sapiens hypothetical protein MGC12900 (MGC12900), mRNA  Homo sapiens hypothetical protein MGC14801 (MGC14801), mRNA
NM 032694	Homo sapiens hypothetical protein MGC12935 (MGC12935), mRNA
	Homo sapiens hypothetical protein MGC12933 (MGC12933), mRNA  Homo sapiens hypothetical protein MGC10646 (MGC10646), mRNA
NM 032693	Homo sapiens hypothetical protein MGC10946 (MGC10946), mRNA  Homo sapiens hypothetical protein MGC10977 (MGC10977), mRNA
NM 032681	Homo sapiens hypothetical protein MGC10977 (MGC10977), mRNA  Homo sapiens hypothetical protein MGC3413 (MGC3413), mRNA
NM_032678	Homo sapiens hypothetical protein MGC3413 (MGC3413), inRNA  Homo sapiens hypothetical protein MGC4694 (MGC4694), mRNA
NM_032667	Homo sapiens hypothetical protein MGC4094 (MGC4094), inRNA  Homo sapiens hypothetical protein MGC5139 (MGC5139), mRNA
NM_032661	Homo sapiens hypothetical protein MGC3139 (MGC3139), mRNA  Homo sapiens hypothetical protein MGC3079 (MGC3079), mRNA
NM_032634	Tromo sapiens hypomenicai protein MGC3073 (MGC3073), mRNA

NM_032631	Homo sapiens hypothetical protein MGC2641 (MGC2641), mRNA
NM_032601	Homo sapiens methylmalonyl CoA epimerase (MCEE), mRNA
NM_032596	Homo sapiens testes development-related NYD-SP22 (NYD-SP22), mRNA
NM_032593	Homo sapiens PKCI-1-related HIT protein (HIT-17), mRNA
NM_032586	Homo sapiens testis transcript Y 8 (TTY8), mRNA
NM_032582	Homo sapiens ubiquitin specific protease (NY-REN-60), mRNA
NM_032580	Homo sapiens hairy and enhancer of split 7 (Drosophila) (HES7), mRNA
NM_032574	Homo sapiens dpy-30-like protein (LOC84661), mRNA
NM_032558	Homo sapiens hypothetical protein FLJ14753 (FLJ14753), mRNA
NM_032557	Homo sapiens HP43.8KD protein (HP43.8KD), mRNA
NM_032553	Homo sapiens putative purinergic receptor (FKSG79), mRNA
NM_032545	Homo sapiens cryptic gene (CRYPTIC), mRNA
NM_020963	Homo sapiens Mov10, Moloney leukemia virus 10, homolog (mouse) (MOV10),
ND 6 022522	mRNA
NM_032522	Homo sapiens hypothetical protein MGC2629 (MGC2629), mRNA
NM_032507	Homo sapiens cerebral protein-4 (HUCEP-4), mRNA
NM_032499	Homo sapiens hypothetical protein HH114 (HH114), mRNA
NM 032494	Homo sapiens zinc finger protein (LOC84524), mRNA
NM_032492	Homo sapiens hypothetical protein GL009 (GL009), mRNA
NM_032487	Homo sapiens actin related protein M1 (ARPM1), mRNA
NM_032486	Homo sapiens dynactin 4 (MGC3248), mRNA
NM_032445	Homo sapiens MEGF11 protein (MEGF11), mRNA
NM_030898	Homo sapiens hypothetical protein FLJ21673 (FLJ21673), mRNA
NM_032412	Homo sapiens putative nuclear protein ORF1-FL49 (ORF1-FL49), mRNA
NM 032411	Homo sapiens esophageal cancer related gene 4 protein (ECRG4), mRNA
NM_015247	Homo sapiens cylindromatosis (turban tumor syndrome) (CYLD), mRNA
NM_032330	Homo sapiens hypothetical protein MGC12536 (MGC12536), mRNA
NM 032384	Homo sapiens hypothetical protein FLJ23183 (FLJ23183), mRNA
NM_032372	Homo sapiens hypothetical protein MGC16186 (MGC16186), mRNA
NM_032367 NM_032354	Homo sapiens hypothetical protein MGC15435 (MGC15435), mRNA
NM 032347	Homo sapiens hypothetical protein MGC10744 (MGC10744), mRNA
NM 032344	Homo sapiens hypothetical protein MGC13250 (MGC13250), mRNA
NM 032342	Homo sapiens hypothetical protein MGC13045 (MGC13045), mRNA
NM 032342 NM 032340	Homo sapiens hypothetical protein MGC12992 (MGC12992), mRNA
NM_032338	Homo sapiens hypothetical protein MGC14833 (MGC14833), mRNA
NM 032333	Homo sapiens hypothetical protein MGC14817 (MGC14817), mRNA
NM 032327	Homo sapiens hypothetical protein MGC4248 (MGC4248), mRNA
NM 032325	Homo sapiens hypothetical protein MGC2993 (MGC2993), mRNA
NM_032324	Homo sapiens hypothetical protein MGC11102 (MGC11102), mRNA
NM 032323	Homo sapiens hypothetical protein MGC13186 (MGC13186), mRNA
NM 032320	Homo sapiens hypothetical protein MGC13102 (MGC13102), mRNA
NM 032318	Homo sapiens hypothetical protein MGC13007 (MGC13007), mRNA
NM_032317	Homo sapiens hypothetical protein MGC12945 (MGC12945), mRNA
NM_032317 NM_032316	Homo sapiens hypothetical protein MGC12943 (MGC12943), mRNA
NM 032305	Homo sapiens hypothetical protein MGC12936 (MGC12936), mRNA
NM_032293	Homo sapiens hypothetical protein MGC3200 (MGC3200), mRNA
NM 032291	Homo sapiens hypothetical protein DKFZp761J1523 (DKFZp761J1523), mRNA
	Homo sapiens hypothetical protein DKFZp761D221 (DKFZp761D221), mRNA
NM_032290	Homo sapiens hypothetical protein DKFZp761C121 (DKFZp761C121), mRNA
NM_032288	Homo sapiens hypothetical protein DKFZp761B1514 (DKFZp761B1514), mRNA
NM_032273	Homo sapiens hypothetical protein DKFZp586C1924 (DKFZp586C1924),
	——————————————————————————————————————

	mRNA PNA
VM 032299	Homo sapiens hypothetical protein MGC2714 (MGC2714), mRNA
NM 032267	Thems genions hymothetical protein DK FZD434E169 (DKFZD434E169), IIKNA
NM 032264	Homo senions hypothetical protein DKFZp434D177 (DKFZp434D177), mRNA
NM_032261	Homo sapiens hypothetical protein DKFZp434N0650 (DKFZp434N0650),
NM_032258	Homo sapiens hypothetical protein DKFZp434P2235 (DKFZp434P2235),
NM_032251	Homo sapiens hypothetical protein DKFZp434G0920 (DKFZp434G0920),
NM 032250	Homo saniens hypothetical protein DKFZp434A171 (DKFZp434A171), mRNA
NM_032249	Homo sapiens hypothetical protein DKFZp434F1819 (DKFZp434F1819),
NM_032248	Homo sapiens hypothetical protein DKFZp434F1719 (DKFZp434F1719),
NM 032246	Homo saniens hypothetical protein DKFZp434J0617 (DKFZp434J0617), mRNA
NM 032245	Homo saniens hypothetical protein DKFZp434I1916 (DKFZp434I1916), mKNA
NM 032223	Homo saniens hypothetical protein FLJ22427 (FLJ22427), mRNA
NM 032209	Homo sapiens hypothetical protein FLJ21777 (FLJ21777), mRNA
NM 032193	Homo sapiens hypothetical protein FLJ20974 (FLJ20974), mRNA
NM 032177	Homo sapiens hypothetical protein FLJ13193 (FLJ13193), mRNA
NM 032167	Homo sapiens hypothetical protein FLJ12363 (FLJ12363), mRNA
NM 032161	Homo sapiens KIAA1870 protein (KIAA1870), mRNA
NM 032154	Homo saniens MRLR protein (MBLR), mRNA
NM_032151	Homo sapiens hypothetical protein DKFZp566K1946 (DKFZP566K1946),
NM_032148	Homo sapiens hypothetical protein DKFZp434K0427 (DKFZP434K0427),
NM_032139	Homo sapiens hypothetical protein DKFZp434L0718 (DKFZP434L0718), mRNA
NM_032138	Homo sapiens hypothetical protein DKFZp434E2318 (DKFZP434E2318),
NM_032136	Homo sapiens hypothetical protein DKFZp434L1717 (DKFZP434L1717),
NM_032125	Homo sapiens hypothetical protein DKFZp564D0478 (DKFZP564D0478),
NM_032120	Homo sapiens hypothetical protein DKFZp564O0523 (DKFZP564O0523), mRNA
NM 020921	Homo saniens ninein (GSK3B interacting protein) (NIN), mRNA
NM 020441	Homo saniens hypothetical protein DKFZp762I166 (DKFZP762I166), mRNA
NM_018719	Homo sapiens hypothetical protein DKFZp762L0311 (DKFZp762L0311), mRNA
NM 015630	Homo sapiens DKFZP566F2124 protein (DKFZP566F2124), mRNA
NM_015621	Homo sapiens DKFZP434C171 protein (DKFZP434C171), mRNA
NM_015595	Homo saniens DKFZP434D146 protein (DKFZP434D146), mRNA
NM_015496	Homo sapiens DKFZP434I116 protein (DKFZP434I116), mRNA
NM 015471	Homo sapiens DKFZP566O1646 protein (DC8), mRNA
NM 015453	Homo sapiens DKFZP434F091 protein (DKFZP434F091), mRNA
NM_015023	Homo sapiens KIAA1037 protein (KIAA1037), mRNA
NM_014972	Homo saniens KIAA 1049 protein (KIAA 1049), mRNA
NM 032042	Homo sapiens hypothetical protein DKFZp564D172 (DKFZP564D172), mRN
NM 032036	Homo sapiens TLH29 protein precursor (TLH29), mRNA

NM_032030 Homo sapiens FKSG83 (FKSG83), mRNA NM_032028 Homo sapiens serine/threonine kinase FKSG81 (FKSG81), mRN NM_032025 Homo sapiens CDA02 protein (CDA02), mRNA NM_032021 Homo sapiens AD031 protein (AD031), mRNA NM_031944 Homo sapiens Mix-like homeobox protein 1 (MILD1), mRNA NM_031920 Homo sapiens ARG99 protein (ARG99), mRNA NM_031480 Homo sapiens hypothetical protein AD034 (AD034), mRNA NM_031477 Homo sapiens hypothetical protein DKFZp434I2117 (DKFZP434 NM_031476 Homo sapiens hypothetical protein MGC10500 (MGC10500), m NM_031471 Homo sapiens hypothetical protein MGC10500 (MGC10500), m NM_031472 Homo sapiens hypothetical protein MGC10966 (MGC10966), m NM_031471 Homo sapiens hypothetical protein MGC10966 (MGC10966), m NM_031472 Homo sapiens hypothetical protein MGC10966 (MGC10966), m NM_031473 Homo sapiens hypothetical protein MGC10966 (MGC10966), m NM_031474 Homo sapiens hypothetical protein MGC4607 (MGC4607), mRN NM_031450 Homo sapiens hypothetical protein MGC4607 (MGC4607), mRN NM_031438 Homo sapiens hypothetical protein MGC4607 (MGC4607), mRN NM_031438 Homo sapiens hypothetical protein MGC5442 (MGC5442), mRN NM_031438 Homo sapiens hypothetical protein MGC5442 (MGC5442), mRN NM_031439 Homo sapiens bypothetical protein DKFZp7611172 (DKFZP761) NM_031497 Homo sapiens bypothetical protein DKFZp564G2022), m NM_031290 Homo sapiens hypothetical protein DKFZp564B1023 (DKFZP564G2022), mRNA NM_031291 Homo sapiens hypothetical protein DKFZp564B1023 (DKFZP564G2022), mRNA NM_031290 Homo sapiens hypothetical protein DKFZp434K1172 (DKFZP440)	4I2117), mRNA RNA 4B044), mRNA RNA RNA mber 8B VA I172), mRNA VA VA
NM 032025 Homo sapiens CDA02 protein (CDA02), mRNA  NM 032021 Homo sapiens AD031 protein (AD031), mRNA  NM 031944 Homo sapiens Mix-like homeobox protein 1 (MILD1), mRNA  NM 031920 Homo sapiens ARG99 protein (ARG99), mRNA  NM 031480 Homo sapiens hypothetical protein AD034 (AD034), mRNA  NM 031478 Homo sapiens hypothetical protein DKFZp434I2117 (DKFZP434-  NM 031477 Homo sapiens hypothetical protein MGC10500 (MGC10500), m  NM 031476 Homo sapiens hypothetical protein DKFZp434B044 (DKFZP434-  NM 031472 Homo sapiens hypothetical protein MGC11134 (MGC11134), m  NM 031471 Homo sapiens hypothetical protein MGC10966 (MGC10966), m  NM 031457 Homo sapiens membrane-spanning 4-domains, subfamily A, mer  (MS4A8B), mRNA  NM 031443 Homo sapiens hypothetical protein MGC4607 (MGC4607), mRN  NM 031438 Homo sapiens hypothetical protein MGC4607 (MGC4607), mRN  NM 031438 Homo sapiens hypothetical protein MGC5442 (MGC5442), mRN  NM 031439 Homo sapiens chromosome 11 open reading frame 25 (C11orf25-  NM 015497 Homo sapiens DKFZP564G2022 protein (DKFZP564G2022), m  NM 031295 Homo sapiens hypothetical protein DKFZp564B1023 (DKFZP56-  mRNA  NM 031291 Homo sapiens hypothetical protein DKFZp434N1235 (DKFZP44-  mRNA	4I2117), mRNA RNA 4B044), mRNA RNA RNA mber 8B  VA I172), mRNA VA O), mRNA
NM_032021 Homo sapiens AD031 protein (AD031), mRNA  NM_031944 Homo sapiens Mix-like homeobox protein 1 (MILD1), mRNA  NM_031920 Homo sapiens ARG99 protein (ARG99), mRNA  NM_031480 Homo sapiens hypothetical protein AD034 (AD034), mRNA  NM_031478 Homo sapiens hypothetical protein DKFZp434I2117 (DKFZP434-  NM_031477 Homo sapiens hypothetical protein MGC10500 (MGC10500), m  NM_031476 Homo sapiens hypothetical protein DKFZp434B044 (DKFZP434-  NM_031472 Homo sapiens hypothetical protein MGC11134 (MGC11134), m  NM_031471 Homo sapiens hypothetical protein MGC10966 (MGC10966), m  NM_031457 Homo sapiens membrane-spanning 4-domains, subfamily A, mer  (MS4A8B), mRNA  NM_031443 Homo sapiens hypothetical protein MGC4607 (MGC4607), mRN  NM_031438 Homo sapiens hypothetical protein MGC4607 (MGC4607), mRN  NM_031434 Homo sapiens hypothetical protein MGC5442 (MGC5442), mRN  NM_031435 Homo sapiens chromosome 11 open reading frame 25 (C11orf25-  NM_015497 Homo sapiens DKFZP564G2022 protein (DKFZP564G2022), m  NM_031306 Homo sapiens hypothetical protein DKFZp564B1023 (DKFZP56-  mRNA  NM_031295 Homo sapiens hypothetical protein DKFZp564B1023 (DKFZP56-  mRNA  NM_031291 Homo sapiens hypothetical protein DKFZp434N1235 (DKFZP44-  mRNA	RNA HB044), mRNA RNA RNA mber 8B NA H172), mRNA NA NA NA NA RNA
NM 031944 Homo sapiens Mix-like homeobox protein 1 (MILD1), mRNA NM 031920 Homo sapiens ARG99 protein (ARG99), mRNA NM 031480 Homo sapiens hypothetical protein AD034 (AD034), mRNA NM 031478 Homo sapiens hypothetical protein DKFZp434I2117 (DKFZP43- NM 031477 Homo sapiens hypothetical protein MGC10500 (MGC10500), m NM 031476 Homo sapiens hypothetical protein DKFZp434B044 (DKFZP434- NM 031472 Homo sapiens hypothetical protein MGC11134 (MGC11134), m NM 031471 Homo sapiens hypothetical protein MGC10966 (MGC10966), m NM 031457 Homo sapiens membrane-spanning 4-domains, subfamily A, mer (MS4A8B), mRNA NM 031443 Homo sapiens hypothetical protein p5326 (P5326), mRNA NM 031443 Homo sapiens hypothetical protein MGC4607 (MGC4607), mRN NM 031434 Homo sapiens hypothetical protein DKFZp761I172 (DKFZP761I NM 031434 Homo sapiens hypothetical protein MGC5442 (MGC5442), mRN NM 031418 Homo sapiens chromosome 11 open reading frame 25 (C11orf25 NM 015497 Homo sapiens DKFZP564G2022 protein (DKFZP564G2022), m NM 031295 Homo sapiens hypothetical protein DKFZp564B1023 (DKFZP56 mRNA NM 031291 Homo sapiens hypothetical protein DKFZp434N1235 (DKFZP44 mRNA	RNA HB044), mRNA RNA RNA mber 8B NA H172), mRNA NA NA NA NA RNA
NM 031480 Homo sapiens ARG99 protein (ARG99), mRNA NM 031480 Homo sapiens hypothetical protein AD034 (AD034), mRNA NM 031478 Homo sapiens hypothetical protein DKFZp434I2117 (DKFZP43- NM 031477 Homo sapiens hypothetical protein MGC10500 (MGC10500), m NM 031476 Homo sapiens hypothetical protein DKFZp434B044 (DKFZP43- NM 031472 Homo sapiens hypothetical protein MGC11134 (MGC11134), m NM 031471 Homo sapiens hypothetical protein MGC10966 (MGC10966), m NM 031457 Homo sapiens membrane-spanning 4-domains, subfamily A, mer (MS4A8B), mRNA NM 031450 Homo sapiens hypothetical protein p5326 (P5326), mRNA NM 031443 Homo sapiens hypothetical protein MGC4607 (MGC4607), mRN NM 031438 Homo sapiens hypothetical protein DKFZp7611172 (DKFZP761) NM 031434 Homo sapiens hypothetical protein MGC5442 (MGC5442), mRN NM 031418 Homo sapiens chromosome 11 open reading frame 25 (C11orf25 NM 015497 Homo sapiens DKFZP564G2022 protein (DKFZP564G2022), m NM 031306 Homo sapiens hypothetical protein DKFZp564B1023 (DKFZP56 mRNA NM 031291 Homo sapiens hypothetical protein DKFZp434N1235 (DKFZP46 mRNA NM 031291 Homo sapiens hypothetical protein DKFZp434N1235 (DKFZP46 mRNA)	RNA HB044), mRNA RNA RNA mber 8B NA H172), mRNA NA NA NA NA RNA
NM 031480 Homo sapiens hypothetical protein AD034 (AD034), mRNA NM 031478 Homo sapiens hypothetical protein DKFZp434I2117 (DKFZP434- NM 031477 Homo sapiens hypothetical protein MGC10500 (MGC10500), m NM 031476 Homo sapiens hypothetical protein DKFZp434B044 (DKFZP434- NM 031472 Homo sapiens hypothetical protein MGC11134 (MGC11134), m NM 031471 Homo sapiens hypothetical protein MGC10966 (MGC10966), m NM 031457 Homo sapiens membrane-spanning 4-domains, subfamily A, mer (MS4A8B), mRNA  NM 031450 Homo sapiens hypothetical protein p5326 (P5326), mRNA NM 031443 Homo sapiens hypothetical protein MGC4607 (MGC4607), mRN NM 031438 Homo sapiens hypothetical protein DKFZp761I172 (DKFZP761) NM 031434 Homo sapiens hypothetical protein MGC5442 (MGC5442), mRN NM 031438 Homo sapiens chromosome 11 open reading frame 25 (C11orf25) NM 015497 Homo sapiens DKFZP564G2022 protein (DKFZP564G2022), m NM 031306 Homo sapiens hypothetical protein DKFZp564B1023 (DKFZP564MRNA  NM 031291 Homo sapiens hypothetical protein DKFZp434N1235 (DKFZP454 mRNA  NM 031291 Homo sapiens hypothetical protein DKFZp434N1235 (DKFZP454 mRNA	RNA HB044), mRNA RNA RNA mber 8B NA H172), mRNA NA NA NA NA RNA
NM 031478 Homo sapiens hypothetical protein DKFZp434I2117 (DKFZP434IM NM 031477 Homo sapiens hypothetical protein MGC10500 (MGC10500), ml NM 031476 Homo sapiens hypothetical protein DKFZp434B044 (DKFZP434IM NM 031472 Homo sapiens hypothetical protein MGC11134 (MGC11134), ml NM 031471 Homo sapiens hypothetical protein MGC10966 (MGC10966), ml NM 031457 Homo sapiens membrane-spanning 4-domains, subfamily A, mer (MS4A8B), mRNA  NM 031450 Homo sapiens hypothetical protein p5326 (P5326), mRNA  NM 031443 Homo sapiens hypothetical protein MGC4607 (MGC4607), mRN  NM 031438 Homo sapiens hypothetical protein DKFZp761I172 (DKFZP761)  NM 031434 Homo sapiens hypothetical protein MGC5442 (MGC5442), mRN  NM 031438 Homo sapiens chromosome 11 open reading frame 25 (C11orf25)  NM 015497 Homo sapiens DKFZP564G2022 protein (DKFZP564G2022), ml NM 031306 Homo sapiens hypothetical protein DKFZp564B1023 (DKFZP564M NM 031291 Homo sapiens hypothetical protein DKFZp564B1023 (DKFZP564M NM 031291 Homo sapiens hypothetical protein DKFZp434N1235 (DKFZP454 mRNA)  NM 031291 Homo sapiens hypothetical protein DKFZp434N1235 (DKFZP454 mRNA)	RNA HB044), mRNA RNA RNA mber 8B NA H172), mRNA NA NA NA NA RNA
NM 031477 Homo sapiens hypothetical protein MGC10500 (MGC10500), m NM 031476 Homo sapiens hypothetical protein DKFZp434B044 (DKFZP434 NM 031472 Homo sapiens hypothetical protein MGC11134 (MGC11134), m NM 031471 Homo sapiens hypothetical protein MGC10966 (MGC10966), m NM 031457 Homo sapiens membrane-spanning 4-domains, subfamily A, mer (MS4A8B), mRNA NM 031450 Homo sapiens hypothetical protein p5326 (P5326), mRNA NM 031443 Homo sapiens hypothetical protein MGC4607 (MGC4607), mRN NM 031438 Homo sapiens hypothetical protein DKFZp761I172 (DKFZP761) NM 031434 Homo sapiens hypothetical protein MGC5442 (MGC5442), mRN NM 031418 Homo sapiens chromosome 11 open reading frame 25 (C11orf25 NM 015497 Homo sapiens DKFZP564G2022 protein (DKFZP564G2022), m NM 031306 Homo sapiens hypothetical protein DKFZp564B1023 (DKFZP56 mRNA NM 031291 Homo sapiens hypothetical protein DKFZp434N1235 (DKFZP42 mRNA	RNA HB044), mRNA RNA RNA mber 8B NA H172), mRNA NA NA NA NA RNA
NM_031476 Homo sapiens hypothetical protein DKFZp434B044 (DKFZP434NM_031472 Homo sapiens hypothetical protein MGC11134 (MGC11134), mNM_031471 Homo sapiens hypothetical protein MGC10966 (MGC10966), mNM_031457 Homo sapiens membrane-spanning 4-domains, subfamily A, memory (MS4A8B), mRNA  NM_031450 Homo sapiens hypothetical protein p5326 (P5326), mRNA  NM_031443 Homo sapiens hypothetical protein MGC4607 (MGC4607), mRN  NM_031438 Homo sapiens hypothetical protein DKFZp761I172 (DKFZP761)  NM_031434 Homo sapiens hypothetical protein MGC5442 (MGC5442), mRN  NM_031418 Homo sapiens chromosome 11 open reading frame 25 (C11orf25)  NM_015497 Homo sapiens DKFZP564G2022 protein (DKFZP564G2022), mRNA  NM_031306 Homo sapiens hypothetical protein DKFZp564B1023 (DKFZP564MRNA)  NM_031291 Homo sapiens hypothetical protein DKFZp434N1235 (DKFZP42)  MRNA	AB044), mRNA RNA RNA mber 8B  NA I172), mRNA NA NA NA NA RNA
NM 031472 Homo sapiens hypothetical protein MGC11134 (MGC11134), m NM 031471 Homo sapiens hypothetical protein MGC10966 (MGC10966), m NM 031457 Homo sapiens membrane-spanning 4-domains, subfamily A, mer (MS4A8B), mRNA  NM 031440 Homo sapiens hypothetical protein p5326 (P5326), mRNA NM 031443 Homo sapiens hypothetical protein MGC4607 (MGC4607), mRN NM 031438 Homo sapiens hypothetical protein DKFZp761I172 (DKFZP761) NM 031434 Homo sapiens hypothetical protein MGC5442 (MGC5442), mRN NM 031418 Homo sapiens chromosome 11 open reading frame 25 (C11orf25) NM 015497 Homo sapiens DKFZP564G2022 protein (DKFZP564G2022), m NM 031306 Homo sapiens hypothetical protein DKFZp564B1023 (DKFZP56 mRNA  NM 031295 Homo sapiens hypothetical protein PP1226 (PP1226), mRNA NM 031291 Homo sapiens hypothetical protein DKFZp434N1235 (DKFZP42) mRNA	RNA RNA mber 8B  VA I172), mRNA VA VA NA NA RNA
NM_031471 Homo sapiens hypothetical protein MGC10966 (MGC10966), m NM_031457 Homo sapiens membrane-spanning 4-domains, subfamily A, mer (MS4A8B), mRNA  NM_031450 Homo sapiens hypothetical protein p5326 (P5326), mRNA NM_031443 Homo sapiens hypothetical protein MGC4607 (MGC4607), mRN NM_031438 Homo sapiens hypothetical protein DKFZp761I172 (DKFZP761) NM_031434 Homo sapiens hypothetical protein MGC5442 (MGC5442), mRN NM_031418 Homo sapiens chromosome 11 open reading frame 25 (C11orf25) NM_015497 Homo sapiens DKFZP564G2022 protein (DKFZP564G2022), m NM_031306 Homo sapiens hypothetical protein DKFZp564B1023 (DKFZP56 mRNA  NM_031295 Homo sapiens hypothetical protein PP1226 (PP1226), mRNA NM_031291 Homo sapiens hypothetical protein DKFZp434N1235 (DKFZP42) mRNA	RNA mber 8B  NA I172), mRNA NA NA N, mRNA RNA
NM_031457 Homo sapiens membrane-spanning 4-domains, subfamily A, mem (MS4A8B), mRNA  NM_031450 Homo sapiens hypothetical protein p5326 (P5326), mRNA  NM_031443 Homo sapiens hypothetical protein MGC4607 (MGC4607), mRN  NM_031438 Homo sapiens hypothetical protein DKFZp761I172 (DKFZP761I  NM_031434 Homo sapiens hypothetical protein MGC5442 (MGC5442), mRN  NM_031418 Homo sapiens chromosome 11 open reading frame 25 (C11orf25  NM_015497 Homo sapiens DKFZP564G2022 protein (DKFZP564G2022), m  NM_031306 Homo sapiens hypothetical protein DKFZp564B1023 (DKFZP56  mRNA  NM_031295 Homo sapiens hypothetical protein PP1226 (PP1226), mRNA  NM_031291 Homo sapiens hypothetical protein DKFZp434N1235 (DKFZP43  mRNA	NA I172), mRNA NA ), mRNA RNA
(MS4A8B), mRNA  NM_031450 Homo sapiens hypothetical protein p5326 (P5326), mRNA  NM_031443 Homo sapiens hypothetical protein MGC4607 (MGC4607), mRN  NM_031438 Homo sapiens hypothetical protein DKFZp761I172 (DKFZP761I  NM_031434 Homo sapiens hypothetical protein MGC5442 (MGC5442), mRN  NM_031418 Homo sapiens chromosome 11 open reading frame 25 (C11orf25  NM_015497 Homo sapiens DKFZP564G2022 protein (DKFZP564G2022), m  NM_031306 Homo sapiens hypothetical protein DKFZp564B1023 (DKFZP56  mRNA  NM_031295 Homo sapiens hypothetical protein PP1226 (PP1226), mRNA  NM_031291 Homo sapiens hypothetical protein DKFZp434N1235 (DKFZP43  mRNA	NA I172), mRNA NA ), mRNA RNA
NM 031450 Homo sapiens hypothetical protein p5326 (P5326), mRNA  NM 031443 Homo sapiens hypothetical protein MGC4607 (MGC4607), mRN  NM 031438 Homo sapiens hypothetical protein DKFZp761I172 (DKFZP761  NM 031434 Homo sapiens hypothetical protein MGC5442 (MGC5442), mRN  NM 031418 Homo sapiens chromosome 11 open reading frame 25 (C11orf25  NM 015497 Homo sapiens DKFZP564G2022 protein (DKFZP564G2022), m  NM 031306 Homo sapiens hypothetical protein DKFZp564B1023 (DKFZP56  mRNA  NM 031295 Homo sapiens hypothetical protein PP1226 (PP1226), mRNA  NM 031291 Homo sapiens hypothetical protein DKFZp434N1235 (DKFZP42  mRNA	I172), mRNA NA ), mRNA RNA
NM 031443 Homo sapiens hypothetical protein MGC4607 (MGC4607), mRN NM 031438 Homo sapiens hypothetical protein DKFZp761I172 (DKFZP761INM 031434 Homo sapiens hypothetical protein MGC5442 (MGC5442), mRN NM 031418 Homo sapiens chromosome 11 open reading frame 25 (C11orf25 NM 015497 Homo sapiens DKFZP564G2022 protein (DKFZP564G2022), mRN 031306 Homo sapiens hypothetical protein DKFZp564B1023 (DKFZP56 mRNA NM 031295 Homo sapiens hypothetical protein PP1226 (PP1226), mRNA NM 031291 Homo sapiens hypothetical protein DKFZp434N1235 (DKFZP42 mRNA)	I172), mRNA NA ), mRNA RNA
NM 031438 Homo sapiens hypothetical protein DKFZp761I172 (DKFZP761INM 031434 Homo sapiens hypothetical protein MGC5442 (MGC5442), mRN NM 031418 Homo sapiens chromosome 11 open reading frame 25 (C11orf25 NM 015497 Homo sapiens DKFZP564G2022 protein (DKFZP564G2022), mNM 031306 Homo sapiens hypothetical protein DKFZp564B1023 (DKFZP56 mRNA NM 031295 Homo sapiens hypothetical protein PP1226 (PP1226), mRNA NM 031291 Homo sapiens hypothetical protein DKFZp434N1235 (DKFZP43 mRNA	I172), mRNA NA ), mRNA RNA
NM 031434 Homo sapiens hypothetical protein MGC5442 (MGC5442), mRN NM 031418 Homo sapiens chromosome 11 open reading frame 25 (C11orf25 NM 015497 Homo sapiens DKFZP564G2022 protein (DKFZP564G2022), m NM 031306 Homo sapiens hypothetical protein DKFZp564B1023 (DKFZP56 mRNA NM 031295 Homo sapiens hypothetical protein PP1226 (PP1226), mRNA NM 031291 Homo sapiens hypothetical protein DKFZp434N1235 (DKFZP43 mRNA	NA ), mRNA RNA
NM_031418 Homo sapiens chromosome 11 open reading frame 25 (C11orf25 NM_015497 Homo sapiens DKFZP564G2022 protein (DKFZP564G2022), m NM_031306 Homo sapiens hypothetical protein DKFZp564B1023 (DKFZP56 mRNA NM_031295 Homo sapiens hypothetical protein PP1226 (PP1226), mRNA NM_031291 Homo sapiens hypothetical protein DKFZp434N1235 (DKFZP43 mRNA	), mRNA RNA
NM_015497 Homo sapiens DKFZP564G2022 protein (DKFZP564G2022), m NM_031306 Homo sapiens hypothetical protein DKFZp564B1023 (DKFZP56 mRNA  NM_031295 Homo sapiens hypothetical protein PP1226 (PP1226), mRNA  NM_031291 Homo sapiens hypothetical protein DKFZp434N1235 (DKFZP42 mRNA	RNA
NM_031306 Homo sapiens hypothetical protein DKFZp564B1023 (DKFZP56 mRNA  NM_031295 Homo sapiens hypothetical protein PP1226 (PP1226), mRNA  NM_031291 Homo sapiens hypothetical protein DKFZp434N1235 (DKFZP43 mRNA	
mRNA NM_031295 Homo sapiens hypothetical protein PP1226 (PP1226), mRNA NM_031291 Homo sapiens hypothetical protein DKFZp434N1235 (DKFZP43 mRNA	54B1023),
NM_031291 Homo sapiens hypothetical protein DKFZp434N1235 (DKFZP43 mRNA	
NM_031291 Homo sapiens hypothetical protein DKFZp434N1235 (DKFZP43mRNA	
	34N1235),
mRNA	34K1172),
NM_031270 Homo sapiens PRO1596 protein (PRO1596), mRNA	
NM_031268 Homo sapiens PRO0461 protein (PRO0461), mRNA	
NM_031217 Homo sapiens hypothetical protein DKFZp434G2226 (DKFZP4: mRNA	34G2226),
NM_013358 Homo sapiens peptidylarginine deiminase type I (hPAD-colony1	0) mRNA
NM_030980 Homo sapiens hypothetical protein FLJ12671 (FLJ12671), mRN	
NM_030954 Homo sapiens hypothetical protein DKFZp564A022 (DKFZP564	
NM_030953 Homo sapiens hypothetical protein DKFZp761E2110 (DKFZP76	
MRNA NM_030941 Homo sapiens exonuclease NEF-sp (LOC81691), mRNA	
NM_030939 Homo sapiens exonuclease NEF-sp (LOC81691), mRNA  NM_030939 Homo sapiens hypothetical protein FLJ12619 (FLJ12619), mRN	Α
NM_030938 Homo sapiens likely ortholog of rat vacuole membrane protein 1	
mRNA NM 020022 Home conjugate display and homeles 2 (Drescabile) (DIARIE)	-DNIA
NM 030932 Homo sapiens diaphanous homolog 3 (Drosophila) (DIAPH3), n	
NM 030927 Homo sapiens hypothetical protein MGC11352 (MGC11352), m	
NM 030925 Homo sapiens hypothetical protein FLJ12577 (FLJ12577), mRN	A
NM 030918 Homo sapiens hypothetical protein My014 (MY014), mRNA	
NM 030911 Homo sapiens protein kinase NYD-SP15 (NYD-SP15), mRNA	· A
NM 030899 Homo sapiens hypothetical protein FLJ23407 (FLJ23407), mRN	A
NM_018657 Homo sapiens myoneurin (MYNN), mRNA	. DNI A
NM 030818 Homo sapiens hypothetical protein MGC10471 (MGC10471), m	
NM_030813 Homo sapiens suppressor of potassium transport defect 3 (SKD)	
NM_030808 Homo sapiens LIS1-interacting protein NUDEL; endooligopeptic (NUDEL), mRNA	
NM_030805   Homo sapiens hypothetical protein DKFZp564L2423 (DKFZP56	64L2423),

	nRNA DNA
	GERD in duced protein (LOCS1558) MKNA
VM_030802	Homo sapiens C/EBP-induced protein (EOCO1556), Marie 1996. Homo sapiens hypothetical protein DKFZp564O1664 (DKFZP564O1664),
<b>→</b>	7374
T ( 020700	Home seniers hypothetical protein AF140225 (AF140225), mRNA
7.5 020702	Home geniens hypothetical protein SP329 (SP329), IIIKNA
T ( 020702	Homo capiens hymothetical protein PP1665 (PP1665), inkina
	xx foliate transporter/carrier (LOC81034), IMRNA
5 5 500 674	II conjone colute carrier family 38, member 1 (SLC36A1), IIIC 71
NM 030672	Home gapiens hypothetical protein FLJ10312 (FLJ10312), HIKNA
NM 024947	xx assigns hymothetical protein FL 112/29 (FLJ12/29), IIIXIA
	that is a protein FI 11 46/(FI 1) 40/), IIIXIVA
NM_024963	Homo sapiens hypothetical protein DKFZp434M0331 (DKFZp434M0331),
NM_017600	mRNA
NT 6 020(52	NG2 motoin (NG3) mRNA
NM_030652	Homo sapiens NG3 protein (NG3), media; Homo sapiens chromosome 6 open reading frame 31 (C6orf31), mRNA
NM_030651	TIions VIA A 1101 protein (KIAA 1191), MKNA
NM_020444	Homo sapiens kirali protein (kirali 1975), mRNA Homo sapiens hypothetical protein MGC5499 (MGC5499), mRNA
NM_024055	Homo sapiens KIAA0810 protein (KIAA0810), mRNA
NM_025154	Homo sapiens KIAA0810 protein (KIMA0810);
NM_017515	Homo sapiens novel protein (HSNOV1), mRNA  Homo sapiens hypothetical protein FLJ12985 (FLJ12985), mRNA
NM_024924	Homo sapiens hypothetical protein PL312903 (PL312303), the Homo sapiens cytochrome b5 outer mitochondrial membrane precursor (CYB5-
NM_030579	
	M), mRNA
NM_022068	Homo sapiens hypothetical protein FLJ23403 (FLJ23403), mRNA
NM_025179	Homo sapiens plexin A2 (PLXNA2), mRNA  Homo sapiens plexin A2 (PLXNA2), mRNA
NM_014033	Homo sapiens DKFZP586A0522 protein (DKFZP586A0522), mRNA Homo sapiens DKFZP586A0522 protein (DKFZP586A0522), mRNA
NM_006468	Homo sapiens DKFZF386A0322 protein (DKFZF386A0322 protein (DKFZF386A032 protein (D
NM_025263	Homo sapiens CAT56 protein (CAT56), mRNA
NM_025262	Homo sapiens G5C protein (G5C), mRNA
NM_025261	Homo sapiens G6C protein (G6C), mRNA
NM 025260	Homo sapiens G6B protein (G6B), mRNA
NM 025259	Homo sapiens NG23 protein (NG23), mRNA
NM 025258	TI MKNA
NM 025231	Homo sapiens hypothetical protein (57-6), FLJ22191 (FLJ22191), mRNA
NM 025226	TI Assigns MCTD032 protein (MSTP032), MKNA
NM 025211	Trans conjens protein kingse anchoring protein GRAP42 (GRAI 42), mid 11
NM 025201	Homo seniens hypothetical protein PP1628 (PP1628), mRNA
NM_025192	Homo saniens hypothetical protein FLJ230/1 (FLJ230/1), HIKNA
NM 025188	Homo saniens hypothetical protein FLJ13181 (FLJ13181), mRNA
NM_025174	Homo saniens hypothetical protein FLJ23040 (FLJ23040), HIKNA
NM 025165	Homo saniens hypothetical protein FLJ22637 (FLJ22637), HIKINA
NM_025160	Homo seniens hypothetical protein FLJ21016 (FLJ21016), IIIKNA
NM 025153	Home seniers hypothetical protein FLJ214// (FLJ214//), mkNA
NM 025151	Homo capiens hypothetical protein FLJ22622 (FLJ22622), HIKNA
NM 025149	Homo capiens hypothetical protein FLJ20920 (FLJ20920), IIIXIX
NM 025144	Homo saniens hypothetical protein FLJ226/0 (FLJ226/0), IIIKNA
NM 025138	Homo sapiens hypothetical protein FLJ12661 (FLJ12661), mRNA
	Home geniens ring finger protein 34 (RNF34), MRNA
NM_025126	Homo saniens hypothetical protein FLJ13263 (FLJ13263), mRNA
NM_025125	Home seniers hymothetical protein FLJ21/49 (FLJ21/49), IIIKNA
NM 025124	
NM 025109	Homo sapiens hypothetical protein FLJ22170 (FLJ22170), mRNA

77.5 005000	
NM_025098	Homo sapiens hypothetical protein FLJ22644 (FLJ22644), mRNA
NM_025097	Homo sapiens hypothetical protein FLJ21106 (FLJ21106), mRNA
NM_025095	Homo sapiens hypothetical protein FLJ23558 (FLJ23558), mRNA
NM_025086	Homo sapiens hypothetical protein FLJ22596 (FLJ22596), mRNA
NM_025080	Homo sapiens hypothetical protein FLJ22316 (FLJ22316), mRNA
NM_025079	Homo sapiens hypothetical protein FLJ23231 (FLJ23231), mRNA
NM_025077	Homo sapiens hypothetical protein FLJ13949 (FLJ13949), mRNA
NM_025076	Homo sapiens hypothetical protein FLJ23591 (FLJ23591), mRNA
NM_025072	Homo sapiens chromosome 9 open reading frame 15 (C9orf15), mRNA
NM_025070	Homo sapiens hypothetical protein FLJ22242 (FLJ22242), mRNA
NM_025058	Homo sapiens hypothetical protein FLJ23229 (FLJ23229), mRNA
NM_025055	Homo sapiens hypothetical protein FLJ23168 (FLJ23168), mRNA
NM_025044	Homo sapiens hypothetical protein FLJ22476 (FLJ22476), mRNA
NM_025043	Homo sapiens hypothetical protein FLJ22404 (FLJ22404), mRNA
NM_025041	Homo sapiens hypothetical protein FLJ22173 (FLJ22173), mRNA
NM_025034	Homo sapiens hypothetical protein FLJ21290 (FLJ21290), mRNA
NM_025032	Homo sapiens hypothetical protein FLJ21272 (FLJ21272), mRNA
NM_025029	Homo sapiens hypothetical protein FLJ14346 (FLJ14346), mRNA
NM_025005	Homo sapiens hypothetical protein FLJ13315 (FLJ13315), mRNA
NM_024998	Homo sapiens hypothetical protein FLJ12704 (FLJ12704), mRNA
NM_024994	Homo sapiens hypothetical protein FLJ12595 (FLJ12595), mRNA
NM_024977	Homo sapiens hypothetical protein FLJ12078 (FLJ12078), mRNA
NM_024976	Homo sapiens hypothetical protein FLJ11996 (FLJ11996), mRNA
NM_024956	Homo sapiens hypothetical protein FLJ23375 (FLJ23375), mRNA
NM_024944	Homo sapiens chromosome 21 open reading frame 68 (C21orf68), mRNA
NM_024942	Homo sapiens hypothetical protein FLJ13490 (FLJ13490), mRNA
NM_024941	Homo sapiens hypothetical protein FLJ13611 (FLJ13611), mRNA
NM_024938	Homo sapiens hypothetical protein FLJ11383 (FLJ11383), mRNA
NM_024935	Homo sapiens hypothetical protein FLJ13687 (FLJ13687), mRNA
NM_024920	Homo sapiens hypothetical protein FLJ14281 (FLJ14281), mRNA
NM_024919	Homo sapiens hypothetical protein FLJ22615 (FLJ22615), mRNA
NM_024917	Homo sapiens hypothetical protein FLJ12687 (FLJ12687), mRNA
NM_024914	Homo sapiens hypothetical protein FLJ13262 (FLJ13262), mRNA
NM_024911	Homo sapiens hypothetical protein FLJ23091 (FLJ23091), mRNA
NM_024909	Homo sapiens hypothetical protein FLJ13158 (FLJ13158), mRNA
NM_024908	Homo sapiens hypothetical protein FLJ12973 (FLJ12973), mRNA
NM_024906	Homo sapiens hypothetical protein FLJ21032 (FLJ21032), mRNA
NM_024897	Homo sapiens hypothetical protein FLJ22672 (FLJ22672), mRNA
NM_024889	Homo sapiens hypothetical protein FLJ23537 (FLJ23537), mRNA
NM_024886	Homo sapiens hypothetical protein FLJ14280 (FLJ14280), mRNA
NM_024882	Homo sapiens hypothetical protein FLJ13189 (FLJ13189), mRNA
NM_024880	Homo sapiens hypothetical protein FLJ23556 (FLJ23556), mRNA
NM_024864	Homo sapiens hypothetical protein FLJ22578 (FLJ22578), mRNA
NM_024853	Homo sapiens hypothetical protein FLJ13385 (FLJ13385), mRNA
NM_024848	Homo sapiens hypothetical protein FLJ13941 (FLJ13941), mRNA
NM_024847	Homo sapiens hypothetical protein FLJ21240 (FLJ21240), mRNA
NM_024841	Homo sapiens hypothetical protein FLJ14213 (FLJ14213), mRNA
NM_024839	Homo sapiens hypothetical protein FLJ22638 (FLJ22638), mRNA
NM_024837	Homo sapiens hypothetical protein FLJ21472 (FLJ21472), mRNA
NM_024835	Homo sapiens C3HC4-type zinc finger protein (LZK1), mRNA
NM_024815	Homo sapiens hypothetical protein FLJ22494 (FLJ22494), mRNA
	, (2202 157), muya

WO 03/074654

NM_024813	Homo sapiens hypothetical protein FLJ13150 (FLJ13150), mRNA
NM_024811	Homo sapiens hypothetical protein FLJ12529 (FLJ12529), mRNA  Homo sapiens hypothetical protein FLJ12529 (FLJ12529), mRNA
NM_024810	Homo sapiens hypothetical protein FLJ23018 (FLJ23018), mRNA  Homo sapiens hypothetical protein FLJ23018 (FLJ23018), mRNA
NM_024809	Homo sapiens hypothetical protein FLJ12975 (FLJ12975), mRNA
NM_024808	Homo sapiens hypothetical protein FLJ22624 (FLJ22624), mRNA
NM 024807	Homo sapiens hypothetical protein FLJ13693 (FLJ13693), mRNA
NM 024806	Homo sapiens hypothetical protein FLJ23554 (FLJ23554), mRNA
NM 024799	Homo sapiens hypothetical protein FLJ13224 (FLJ13224), mRNA
NM 024796	Homo sapiens hypothetical protein FLJ22639 (FLJ22639), mRNA
NM 024789	Homo caniens hypothetical protein FLJ22529 (FLJ22529), IIIXIVA
NM 024784	Home seniers hypothetical protein FLJ23392 (FLJ23392), IIIRNA
NM_024780	Homo saniens hypothetical protein FLJ13593 (FLJ13593), IIIXNA
NM 024773	Homo saniens hypothetical protein FLJ13/98 (FLJ13/98), IRKNA
NM 024772	Homo sapiens hypothetical protein FLJ23151 (FLJ23151), mRNA
NM 024771	Homo saniens hypothetical protein FLJ13848 (FLJ13848), mkina
NM 024763	Homo sapiens hypothetical protein FLJ23129 (FLJ23129), mRNA
NM 024754	Homo seniens hypothetical protein FLJ12598 (FLJ12598), mRNA
NM 024749	Homo conjens hypothetical protein FLJ12505 (FLJ12505), mRNA
NM 024746	Homo saniens hypothetical protein FLJ13840 (FLJ13840), MRNA
NM 024732	Theme conjens hypothetical protein FL/14351 (FL/14351), IIIKNA
NM 024731	Homo saniens chromosome 16 open reading frame 44 (C1001144), inktyA
NM 024727	Homo sapiens hypothetical protein FLJ23259 (FLJ23259), mRNA
NM 024722	Homo sapiens hypothetical protein FLJ13322 (FLJ13322), mRNA
NM 024717	Homo saniens hypothetical protein FLJ22344 (FLJ22344), IIIKNA
NM 024715	Homo sapiens hypothetical protein FLJ22625 (FLJ22625), mRNA
NM_024709	Homo sapiens hypothetical protein FLJ14146 (FLJ14146), mRNA
NM 024705	Homo sapiens hypothetical protein FLJ13639 (FLJ13639), mRNA  Homo sapiens hypothetical protein FLJ13639 (FLJ13639), mRNA
NM_024703	Homo sapiens hypothetical protein FLJ22593 (FLJ22593), mRNA  Homo sapiens hypothetical protein FLJ22593 (FLJ22593), mRNA
NM_024701	Homo sapiens ankyrin repeat and SOCS box-containing 13 (ASB13), mRNA  Homo sapiens ankyrin repeat and SOCS box-containing 13 (ASB13), mRNA
NM_024700	Homo sapiens Smad nuclear interacting protein (SNIP1), mRNA  Homo sapiens Smad nuclear interacting protein (SNIP1), mRNA
NM_024695	Homo sapiens hypothetical protein FLJ13993 (FLJ13993), mRNA  Homo sapiens hypothetical protein FLJ13993 (FLJ13993), mRNA
NM_024693	Homo sapiens hypothetical protein FLJ20909 (FLJ20909), mRNA  Homo sapiens hypothetical protein FLJ20909 (FLJ20909), mRNA
NM_024688	Homo sapiens hypothetical protein FLJ13031 (FLJ13031), mRNA
NM_024686	Homo sapiens hypothetical protein FLJ23033 (FLJ23033), mRNA  Homo sapiens hypothetical protein FLJ23033 (FLJ23033), mRNA
NM_024678	Homo sapiens hypothetical protein FLJ23441 (FLJ23441), mRNA  Homo sapiens hypothetical protein FLJ23441 (FLJ23441), mRNA
NM_024675	Homo sapiens hypothetical protein FLJ21816 (FLJ21816), mRNA
NM_024672	Homo sapiens hypothetical protein FLJ23320 (FLJ23320), mRNA
NM_024666	Homo sapiens hypothetical protein FLJ11506 (FLJ11506), mRNA
NM_024654	Homo sapiens hypothetical protein FLJ23323 (FLJ23323), mRNA
NM_024650	Homo sapiens hypothetical protein FLJ22531 (FLJ22531), mRNA
NM_024649	Homo sapiens hypothetical protein FLJ23590 (FLJ23590), mRNA
NM_024647	Homo sapiens hypothetical protein FLJ13287 (FLJ13287), mRNA
NM_024640	
NM_024636	Homo sapiens likely ortholog of mouse tumor necrosis-alpha-induced adipose-
	related protein (FLJ23153), mRNA  Homo sapiens hypothetical protein FLJ23188 (FLJ23188), mRNA
NM_024628	
NM_024627	
NM_024626	- T - T - T - T - T - T - T - T - T - T
NM_024624	- 1
NM_024616	
NM_024615	Homo sapiens hypothetical protein i Eszisos (i Eszisos), mad a

NM 024610   Homo sapiens hypothetical protein FLJ22623 (FLJ22623), mRNA   NM 024606   Homo sapiens hypothetical protein FLJ11841 (FLJ21841), mRNA   NM 024606   Homo sapiens hypothetical protein FLJ1120896 (FLJ1120896), mRNA   NM 024602   Homo sapiens hypothetical protein FLJ120896 (FLJ120896), mRNA   NM 024602   Homo sapiens hypothetical protein FLJ12166 (FLJ12166), mRNA   NM 024595   Homo sapiens hypothetical protein FLJ12166 (FLJ12166), mRNA   NM 024595   Homo sapiens hypothetical protein FLJ12166 (FLJ12166), mRNA   NM 024585   Homo sapiens hypothetical protein FLJ13066 (FLJ12666), mRNA   NM 024585   Homo sapiens hypothetical protein FLJ13146 (FLJ131666), mRNA   NM 024580   Homo sapiens hypothetical protein FLJ13119 (FLJ13119), mRNA   NM 024565   Homo sapiens hypothetical protein FLJ13119 (FLJ17112), mRNA   NM 024565   Homo sapiens hypothetical protein FLJ111712 (FLJ171712), mRNA   NM 024565   Homo sapiens hypothetical protein FLJ1103 (FLJ2103), mRNA   NM 024556   Homo sapiens hypothetical protein FLJ121089 (FLJ12089), mRNA   NM 024546   Homo sapiens hypothetical protein FLJ12089 (FLJ12089), mRNA   NM 024546   Homo sapiens hypothetical protein FLJ12089 (FLJ12089), mRNA   NM 024532   Homo sapiens hypothetical protein FLJ12684 (FLJ12684), mRNA   NM 024526   Homo sapiens hypothetical protein FLJ12684 (FLJ12684), mRNA   NM 024526   Homo sapiens hypothetical protein FLJ12680 (FLJ12520), mRNA   NM 024526   Homo sapiens hypothetical protein FLJ12680 (FLJ12520), mRNA   NM 024526   Homo sapiens hypothetical protein FLJ12089 (FLJ12680), mRNA   NM 024526   Homo sapiens hypothetical protein FLJ12089 (FLJ12680), mRNA   NM 024526   Homo sapiens hypothetical protein MGC4666 (MGC4666), mRNA   NM 024516   Homo sapiens hypothetical protein MGC4666 (MGC4666), mRNA   NM 024314   Homo sapiens hypothetical protein MGC4666 (MGC4666), mRNA   NM 024326   Homo sapiens hypothetical protein MGC4666 (MGC4666), mRNA   NM 024326   Homo sapiens hypothetical protein MGC4667 (MGC4679), mRNA   NM 024326   Homo sapiens hypothetical protein MG	ND 6 004612	1.000
NM 024609 Homo sapiens hypothetical protein FLJ21841 (FLJ21841), mRNA NM 024606 Homo sapiens hypothetical protein FLJ210896 (FLJ208996), mRNA NM 024605 Homo sapiens hypothetical protein FLJ20896 (FLJ208996), mRNA NM 024595 Homo sapiens hypothetical protein FLJ2166 (FLJ21156), mRNA NM 024595 Homo sapiens hypothetical protein FLJ22160 (FLJ22160), mRNA NM 024595 Homo sapiens hypothetical protein FLJ22160 (FLJ22160), mRNA NM 024584 Homo sapiens hypothetical protein FLJ3119 (FLJ3119), mRNA NM 024584 Homo sapiens hypothetical protein FLJ3119 (FLJ3119), mRNA NM 024580 Homo sapiens hypothetical protein FLJ31119 (FLJ311712), mRNA NM 024550 Homo sapiens hypothetical protein FLJ31119 (FLJ311712), mRNA NM 024551 Homo sapiens hypothetical protein FLJ3103 (FLJ21103), mRNA NM 024552 Homo sapiens hypothetical protein FLJ3103 (FLJ21103), mRNA NM 024552 Homo sapiens hypothetical protein FLJ3103 (FLJ21103), mRNA NM 024546 Homo sapiens hypothetical protein FLJ31849 (FLJ3449), mRNA NM 024534 Homo sapiens hypothetical protein FLJ31849 (FLJ3449), mRNA NM 024532 Homo sapiens hypothetical protein FLJ21684 (FLJ3684), mRNA NM 024523 Homo sapiens hypothetical protein FLJ25274 (FLJ22724), mRNA NM 024523 Homo sapiens hypothetical protein FLJ21035 (FLJ22035), mRNA NM 024523 Homo sapiens hypothetical protein FLJ3035 (FLJ22035), mRNA NM 024524 Homo sapiens hypothetical protein FLJ3050 (FLJ2050), mRNA NM 024514 Homo sapiens hypothetical protein FLJ3660 (FLJ3650), mRNA NM 024514 Homo sapiens hypothetical protein MGC4606 (MGC4666), mRNA NM 024514 Homo sapiens hypothetical protein MGC4606 (MGC4666), mRNA NM 024514 Homo sapiens hypothetical protein MGC4060 (MGC4606), mRNA NM 024514 Homo sapiens hypothetical protein MGC40791 (MGC10791), mRNA NM 024319 Homo sapiens hypothetical protein MGC4060 (MGC4663), mRNA NM 024310 Homo sapiens hypothetical protein MGC40709 (MGC4179), mRNA NM 024311 Homo sapiens hypothetical protein MGC4070 (MGC4170), mRNA NM 024301 Homo sapiens hypothetical protein MGC4070 (MGC4171), mRNA NM 024301 Homo sapiens hypothetical prot	NM_024613	Homo sapiens phafin 2 (FLJ13187), mRNA
NM 024605 Homo sapiens hypothetical protein FLJ11756 (FLJ11756), mRNA NM 024605 Homo sapiens hypothetical protein FLJ20896 (FLJ20896), mRNA NM 024602 Homo sapiens hypothetical protein FLJ21156 (FLJ21150), mRNA NM 024595 Homo sapiens hypothetical protein FLJ21666 (FLJ12666), mRNA NM 024585 Homo sapiens hypothetical protein FLJ2160 (FLJ22160), mRNA NM 024584 Homo sapiens hypothetical protein FLJ31646 (FLJ13646), mRNA NM 024580 Homo sapiens hypothetical protein FLJ3119 (FLJ13119), mRNA NM 024580 Homo sapiens hypothetical protein FLJ13119 (FLJ13119), mRNA NM 024570 Homo sapiens hypothetical protein FLJ11712 (FLJ11712), mRNA NM 024565 Homo sapiens hypothetical protein FLJ1166 (FLJ14166), mRNA NM 024556 Homo sapiens hypothetical protein FLJ1349 (FLJ13449), mRNA NM 024540 Homo sapiens hypothetical protein FLJ13449 (FLJ13449), mRNA NM 024544 Homo sapiens hypothetical protein FLJ12089), mRNA NM 024545 Homo sapiens hypothetical protein FLJ12224 (FLJ22724), mRNA NM 024546 Homo sapiens hypothetical protein FLJ12252 (FLZ1252), mRNA NM 024526 Homo sapiens hypothetical protein FLJ2252 (FLZ1252), mRNA NM 024526 Homo sapiens hypothetical protein FLJ2252 (FLZ1252), mRNA NM 024526 Homo sapiens hypothetical protein FLJ252 (FLZ1252), mRNA NM 024526 Homo sapiens hypothetical protein FLJ2650 (FLJ2650), mRNA NM 024527 Homo sapiens hypothetical protein MGC4666 (MGC4660), mRNA NM 024516 Homo sapiens hypothetical protein MGC4666 (MGC4660), mRNA NM 024517 Homo sapiens hypothetical protein MGC4663 (MGC4660), mRNA NM 024518 Homo sapiens hypothetical protein MGC4067 (MGC409), mRNA NM 024519 Homo sapiens hypothetical protein MGC4663 (MGC4660), mRNA NM 02430 Homo sapiens hypothetical protein MGC4067 (MGC4179), mRNA NM 02430 Homo sapiens hypothetical protein MGC4067 (MGC4179), mRNA NM 02431 Homo sapiens hypothetical protein MGC4079 (MGC4179), mRNA NM 024321 Homo sapiens hypothetical protein MGC4170 (MGC4179), mRNA NM 02430 Homo sapiens hypothetical protein MGC4170 (MGC4179), mRNA NM 02431 Homo sapiens hypothetical protein MGC4070 (MGC4077), mRN		
NM 024605 Homo sapiens hypothetical protein FLJ21156 (FLJ20896), nRNA NM 024595 Homo sapiens hypothetical protein FLJ21156 (FLJ21156), mRNA NM 024595 Homo sapiens hypothetical protein FLJ12666 (FLJ12666), mRNA NM 024585 Homo sapiens hypothetical protein FLJ12660 (FLJ12660), mRNA NM 024584 Homo sapiens hypothetical protein FLJ1319 (FLJ13119), mRNA NM 024585 Homo sapiens hypothetical protein FLJ13119 (FLJ13119), mRNA NM 024586 Homo sapiens hypothetical protein FLJ13119 (FLJ13119), mRNA NM 024561 Homo sapiens hypothetical protein FLJ11712 (FLJ11712), mRNA NM 024565 Homo sapiens hypothetical protein FLJ1103 (FLJ21103), mRNA NM 024556 Homo sapiens hypothetical protein FLJ1103 (FLJ21103), mRNA NM 024556 Homo sapiens hypothetical protein FLJ12089 (FLJ12089), mRNA NM 024546 Homo sapiens hypothetical protein FLJ13449 (FLJ13449), mRNA NM 024545 Homo sapiens hypothetical protein FLJ12684 (FLJ12684), mRNA NM 024524 Homo sapiens hypothetical protein FLJ22724 (FLJ22724), mRNA NM 024523 Homo sapiens hypothetical protein FLJ22035 (FLJ22035), mRNA NM 024523 Homo sapiens hypothetical protein FLJ21650 (FLJ12650), mRNA NM 024521 Homo sapiens hypothetical protein FLJ2650 (FLJ12650), mRNA NM 024514 Homo sapiens hypothetical protein MGC1660 (MGC4660), mRNA NM 024514 Homo sapiens hypothetical protein MGC10791 (MGC10791), mRNA NM 024514 Homo sapiens hypothetical protein MGC10791 (MGC10791), mRNA NM 024514 Homo sapiens hypothetical protein MGC10791 (MGC10791), mRNA NM 024316 Homo sapiens hypothetical protein MGC10791 (MGC10791), mRNA NM 024317 Homo sapiens hypothetical protein MGC10791 (MGC10791), mRNA NM 024318 Homo sapiens hypothetical protein MGC10791 (MGC10791), mRNA NM 024319 Homo sapiens hypothetical protein MGC10765 (MGC10479), mRNA NM 024310 Homo sapiens hypothetical protein MGC10765 (MGC10479), mRNA NM 024311 Homo sapiens hypothetical protein MGC10765 (MGC10433), mRNA NM 024311 Homo sapiens hypothetical protein MGC3067 (MGC3067), mRNA NM 024301 Homo sapiens hypothetical protein MGC3067 (MGC3171), mRNA NM 024301 Homo sapiens hyp		
NM 024502 Homo sapiens hypothetical protein FLJ21156 (FLJ21150), mRNA NM 024585 Homo sapiens hypothetical protein FLJ12666 (FLJ12666), mRNA NM 024584 Homo sapiens hypothetical protein FLJ2160 (FLJ22160), mRNA NM 024585 Homo sapiens hypothetical protein FLJ31646 (FLJ13640), mRNA NM 024580 Homo sapiens hypothetical protein FLJ3119 (FLJ13119), mRNA NM 024570 Homo sapiens hypothetical protein FLJ11712 (FLJ11712), mRNA NM 024565 Homo sapiens hypothetical protein FLJ17112 (FLJ11712), mRNA NM 024556 Homo sapiens hypothetical protein FLJ1103 (FLJ21103), mRNA NM 024555 Homo sapiens hypothetical protein FLJ1103 (FLJ21103), mRNA NM 024554 Homo sapiens hypothetical protein FLJ12039 (FLJ21039), mRNA NM 024534 Homo sapiens hypothetical protein FLJ12899 (FLJ12089), mRNA NM 024534 Homo sapiens hypothetical protein FLJ12684 (FLJ12684), mRNA NM 024534 Homo sapiens hypothetical protein FLJ22724 (FLJ22724), mRNA NM 024526 Homo sapiens hypothetical protein FLJ22152 (FLJ227224), mRNA NM 024526 Homo sapiens hypothetical protein FLJ22035 (FLJ22035), mRNA NM 024527 Homo sapiens hypothetical protein FLJ2505 (FLJ2035), mRNA NM 024528 Homo sapiens hypothetical protein FLJ2650 (FLJ2650), mRNA NM 024516 Homo sapiens hypothetical protein MGC4666 (MGC4666), mRNA NM 024516 Homo sapiens hypothetical protein MGC4666 (MGC4666), mRNA NM 024516 Homo sapiens hypothetical protein MGC4666 (MGC4666), mRNA NM 024516 Homo sapiens hypothetical protein MGC40791 (MGC10791), mRNA NM 024314 Homo sapiens hypothetical protein MGC4666 (MGC4663), mRNA NM 024314 Homo sapiens hypothetical protein MGC40791 (MGC10791), mRNA NM 024340 Homo sapiens hypothetical protein MGC4079 (MGC4179), mRNA NM 024340 Homo sapiens hypothetical protein MGC40766 (MGC4663), mRNA NM 02431 Homo sapiens hypothetical protein MGC4179 (MGC4179), mRNA NM 02431 Homo sapiens hypothetical protein MGC40766 (MGC406470), mRNA NM 02431 Homo sapiens hypothetical protein MGC4170 (MGC4171), mRNA NM 02431 Homo sapiens hypothetical protein MGC4770 (MGC4170), mRNA NM 02431 Homo sapiens hypothetical protein		
NM 024595 Homo sapiens hypothetical protein FLJ12666 (FLJ12666), mRNA NM 024584 Homo sapiens hypothetical protein FLJ22160 (FLJ22160), mRNA NM 024580 Homo sapiens hypothetical protein FLJ13119 (FLJ131440, mRNA NM 024570 Homo sapiens hypothetical protein FLJ13119 (FLJ13119), mRNA NM 024565 Homo sapiens hypothetical protein FLJ111712 (FLJ11712), mRNA NM 024565 Homo sapiens hypothetical protein FLJ11103 (FLJ1103), mRNA NM 024556 Homo sapiens hypothetical protein FLJ1103 (FLJ21103), mRNA NM 024556 Homo sapiens hypothetical protein FLJ12103 (FLJ21103), mRNA NM 024551 Homo sapiens hypothetical protein FLJ12089 (FLJ12089), mRNA NM 024534 Homo sapiens hypothetical protein FLJ12684 (FLJ12684), mRNA NM 024534 Homo sapiens hypothetical protein FLJ12684 (FLJ12684), mRNA NM 024532 Homo sapiens hypothetical protein FLJ22724 (FLJ22724), mRNA NM 024523 Homo sapiens hypothetical protein FLJ22035 (FLJ22035), mRNA NM 024523 Homo sapiens hypothetical protein FLJ22035 (FLJ22035), mRNA NM 024521 Homo sapiens hypothetical protein MGC4606 (MGC4606), mRNA NM 024522 Homo sapiens hypothetical protein MGC4606 (MGC4606), mRNA NM 024516 Homo sapiens hypothetical protein MGC4603 (MGC4603), mRNA NM 024517 Homo sapiens hypothetical protein MGC10791 (MGC10791), mRNA NM 024507 Homo sapiens hypothetical protein MGC10791 (MGC10791), mRNA NM 024341 Homo sapiens hypothetical protein MGC10791 (MGC10791), mRNA NM 024345 Homo sapiens hypothetical protein MGC10765 (MGC10765), mRNA NM 024340 Homo sapiens hypothetical protein MGC4179 (MGC11779), mRNA NM 024341 Homo sapiens hypothetical protein MGC4179 (MGC11779), mRNA NM 024321 Homo sapiens hypothetical protein MGC4179 (MGC4177), mRNA NM 024321 Homo sapiens hypothetical protein MGC4170 (MGC4177), mRNA NM 02430 Homo sapiens hypothetical protein MGC4170 (MGC4171), mRNA NM 02430 Homo sapiens hypothetical protein MGC4171 (MGC4171), mRNA NM 02431 Homo sapiens hypothetical protein MGC4770 (MGC4777), mRNA NM 02495 Homo sapiens hypothetical protein MGC36506 (MGC3067), mRNA NM 024094 Homo sapiens hypothetical prot		
NM 024585 Homo sapiens hypothetical protein FLJ22160 (FLJ22160), mRNA NM 024584 Homo sapiens hypothetical protein FLJ13646 (FLJ13646), mRNA NM 024580 Homo sapiens hypothetical protein FLJ13119 (FLJ13119), mRNA NM 024570 Homo sapiens hypothetical protein FLJ11712 (FLJ13119), mRNA NM 024565 Homo sapiens hypothetical protein FLJ117103 (FLJ11103), mRNA NM 024556 Homo sapiens hypothetical protein FLJ21103 (FLJ21103), mRNA NM 024552 Homo sapiens hypothetical protein FLJ210389 (FLJ21089), mRNA NM 024554 Homo sapiens hypothetical protein FLJ12689 (FLJ12089), mRNA NM 024534 Homo sapiens hypothetical protein FLJ12684 (FLJ12684), mRNA NM 024534 Homo sapiens hypothetical protein FLJ21524 (FLJ22724), mRNA NM 024532 Homo sapiens hypothetical protein FLJ21522 (FLJ21522), mRNA NM 024523 Homo sapiens hypothetical protein FLJ21525 (FLJ21522), mRNA NM 024526 Homo sapiens hypothetical protein FLJ21650 (FLJ12650), mRNA NM 024516 Homo sapiens hypothetical protein MGC4606 (MGC4606), mRNA NM 024516 Homo sapiens hypothetical protein MGC4606 (MGC4606), mRNA NM 024516 Homo sapiens hypothetical protein MGC4606 (MGC4606), mRNA NM 024518 Homo sapiens hypothetical protein MGC4606 (MGC4606), mRNA NM 024519 Homo sapiens hypothetical protein MGC4606 (MGC4606), mRNA NM 024310 Homo sapiens hypothetical protein MGC10791 (MGC10791), mRNA NM 024310 Homo sapiens hypothetical protein MGC10765 (MGC10765), mRNA NM 024314 Homo sapiens hypothetical protein MGC1179 (MGC1179), mRNA NM 024310 Homo sapiens hypothetical protein MGC1179 (MGC1179), mRNA NM 024310 Homo sapiens hypothetical protein MGC1179 (MGC1179), mRNA NM 024311 Homo sapiens hypothetical protein MGC1179 (MGC4179), mRNA NM 024312 Homo sapiens hypothetical protein MGC10765 (MGC3067), mRNA NM 024314 Homo sapiens hypothetical protein MGC407170 (MGC4179), mRNA NM 024315 Homo sapiens hypothetical protein MGC40707 (MGC4179), mRNA NM 024316 Homo sapiens hypothetical protein MGC10765 (MGC3067), mRNA NM 024311 Homo sapiens hypothetical protein MGC1770 (MGC41770), mRNA NM 024094 Homo sapiens hypothetical		
NM 024584 Homo sapiens hypothetical protein FLJ13646 (FLJ13646), mRNA NM 024570 Homo sapiens hypothetical protein FLJ13119 (FLJ13119), mRNA NM 024565 Homo sapiens hypothetical protein FLJ111712 (FLJ11712), mRNA NM 024565 Homo sapiens hypothetical protein FLJ1166 (FLJ14166), mRNA NM 024556 Homo sapiens hypothetical protein FLJ12103 (FLJ1103), mRNA NM 024556 Homo sapiens hypothetical protein FLJ121089 (FLJ12089), mRNA NM 024546 Homo sapiens hypothetical protein FLJ121089 (FLJ12089), mRNA NM 024544 Homo sapiens hypothetical protein FLJ12684 (FLJ13644), mRNA NM 024534 Homo sapiens hypothetical protein FLJ12684 (FLJ12684), mRNA NM 024525 Homo sapiens hypothetical protein FLJ12684 (FLJ127274), mRNA NM 024526 Homo sapiens hypothetical protein FLJ22727 (FLJ22724), mRNA NM 024527 Homo sapiens hypothetical protein FLJ22035 (FLJ22035), mRNA NM 024528 Homo sapiens hypothetical protein FLJ21650 (FLJ21650), mRNA NM 024516 Homo sapiens hypothetical protein MGC4606 (MGC4666), mRNA NM 024516 Homo sapiens hypothetical protein MGC4606 (MGC4666), mRNA NM 024517 Homo sapiens hypothetical protein MGC10791 (MGC10791), mRNA NM 024319 Homo sapiens Phosphateidal protein MGC10791 (MGC10791), mRNA NM 024340 Homo sapiens Phosphateidal protein MGC10765 (MGC10765), mRNA NM 024330 Homo sapiens hypothetical protein MGC10765 (MGC10765), mRNA NM 024330 Homo sapiens hypothetical protein MGC10765 (MGC10765), mRNA NM 024331 Homo sapiens hypothetical protein MGC10765 (MGC10765), mRNA NM 024332 Homo sapiens hypothetical protein MGC10765 (MGC10765), mRNA NM 024334 Homo sapiens hypothetical protein MGC10765 (MGC4179), mRNA NM 024330 Homo sapiens hypothetical protein MGC10765 (MGC4179), mRNA NM 024311 Homo sapiens hypothetical protein MGC4179 (MGC4179), mRNA NM 024312 Homo sapiens hypothetical protein MGC4170 (MGC4171), mRNA NM 024308 Homo sapiens hypothetical protein MGC4170 (MGC4171), mRNA NM 024091 Homo sapiens hypothetical protein MGC3067 (MGC3067), mRNA NM 024092 Homo sapiens hypothetical protein MGC3067 (MGC3067), mRNA NM 024094 Homo sapiens hypot		
NM 024570 Homo sapiens hypothetical protein FLJ13119 (FLJ13119), mRNA NM 024565 Homo sapiens hypothetical protein FLJ11712 (FLJ11712), mRNA NM 024565 Homo sapiens hypothetical protein FLJ11716 (FLJ14166), mRNA NM 024565 Homo sapiens hypothetical protein FLJ21103 (FLJ21103), mRNA NM 024556 Homo sapiens hypothetical protein FLJ21103 (FLJ21103), mRNA NM 024552 Homo sapiens hypothetical protein FLJ12089 (FLJ12089), mRNA NM 024534 Homo sapiens hypothetical protein FLJ13449 (FLJ13449), mRNA NM 024532 Homo sapiens hypothetical protein FLJ22724 (FLJ227244), mRNA NM 024526 Homo sapiens hypothetical protein FLJ22152 (FLJ21522), mRNA NM 024526 Homo sapiens hypothetical protein FLJ22152 (FLJ21522), mRNA NM 024526 Homo sapiens hypothetical protein FLJ21253 (FLJ21523), mRNA NM 024521 Homo sapiens hypothetical protein FLJ21253 (FLJ21523), mRNA NM 024522 Homo sapiens hypothetical protein MGC4606 (MGC4606), mRNA NM 024516 Homo sapiens hypothetical protein MGC4606 (MGC4606), mRNA NM 024507 Homo sapiens hypothetical protein MGC10791 (MGC10791), mRNA NM 024508 Homo sapiens RIAA0239 protein (KIAA0239), mRNA NM 024419 Homo sapiens Phosphatidylglycerophosphate Synthase (PGS1), mRNA NM 024340 Homo sapiens hypothetical protein MGC10765 (MGC10765), mRNA NM 024340 Homo sapiens hypothetical protein MGC1179 (MGC10765), mRNA NM 024321 Homo sapiens hypothetical protein MGC1179 (MGC1179), mRNA NM 024321 Homo sapiens hypothetical protein MGC1179 (MGC1179), mRNA NM 024321 Homo sapiens hypothetical protein MGC1179 (MGC11779), mRNA NM 024321 Homo sapiens hypothetical protein MGC1179 (MGC1177), mRNA NM 024308 Homo sapiens hypothetical protein MGC1179 (MGC1177), mRNA NM 024308 Homo sapiens hypothetical protein MGC4170 (MGC4170), mRNA NM 024308 Homo sapiens hypothetical protein MGC4170 (MGC4171), mRNA NM 024091 Homo sapiens hypothetical protein MGC3067 (MGC3067), mRNA NM 024114 Homo sapiens hypothetical protein MGC306 (MGC3067), mRNA NM 024092 Homo sapiens hypothetical protein MGC306 (MGC3067), mRNA NM 024094 Homo sapiens hypothetical protein MGC307		
NM 024570 Homo sapiens hypothetical protein FLJ11712 (FLJ11712), mRNA NM 024556 Homo sapiens hypothetical protein FLJ121103 (FLJ21103), mRNA NM 024556 Homo sapiens hypothetical protein FLJ12103 (FLJ21103), mRNA NM 024556 Homo sapiens hypothetical protein FLJ121089 (FLJ12089), mRNA NM 024552 Homo sapiens hypothetical protein FLJ12089 (FLJ12089), mRNA NM 024544 Homo sapiens hypothetical protein FLJ12089 (FLJ13449), mRNA NM 024534 Homo sapiens hypothetical protein FLJ12684 (FLJ13449), mRNA NM 024532 Homo sapiens hypothetical protein FLJ122724 (FLJ22724), mRNA NM 024526 Homo sapiens hypothetical protein FLJ21522 (FLJ21522), mRNA NM 024521 Homo sapiens hypothetical protein FLJ21525 (FLJ21522), mRNA NM 024522 Homo sapiens hypothetical protein FLJ212650 (FLJ12650), mRNA NM 024516 Homo sapiens hypothetical protein FLJ12650 (FLJ12650), mRNA NM 024514 Homo sapiens hypothetical protein MGC4603 (MGC4663), mRNA NM 024515 Homo sapiens hypothetical protein MGC4603 (MGC4663), mRNA NM 024516 Homo sapiens hypothetical protein MGC10791 (MGC10791), mRNA NM 024517 Homo sapiens KIAA0239 protein (KIAA0239), mRNA NM 024319 Homo sapiens Phosphatidylglycerophosphate Synthase (PGS1), mRNA NM 024340 Homo sapiens hypothetical protein MGC10765 (MGC10765), mRNA NM 024340 Homo sapiens hypothetical protein MGC4170 (MGC4179), mRNA NM 024321 Homo sapiens hypothetical protein MGC4170 (MGC4179), mRNA NM 024312 Homo sapiens hypothetical protein MGC10743 (MGC10433), mRNA NM 024314 Homo sapiens hypothetical protein MGC10433 (MGC10433), mRNA NM 024316 Homo sapiens hypothetical protein MGC10470 (MGC4170), mRNA NM 024317 Homo sapiens hypothetical protein MGC10433 (MGC10433), mRNA NM 024318 Homo sapiens hypothetical protein MGC10470 (MGC4170), mRNA NM 024319 Homo sapiens hypothetical protein MGC10470 (MGC4170), mRNA NM 024311 Homo sapiens hypothetical protein MGC3067 (MGC3067), mRNA NM 024091 Homo sapiens hypothetical protein MGC3067 (MGC3067), mRNA NM 024092 Homo sapiens hypothetical protein MGC3067 (MGC3067), mRNA NM 024094 Homo sapiens hypothetical pr		
NM 024555 Homo sapiens hypothetical protein FLJ14166 (FLJ14166), mRNA NM 024556 Homo sapiens hypothetical protein FLJ21103 (FLJ21103), mRNA NM 024552 Homo sapiens hypothetical protein FLJ21089 (FLJ12089), mRNA NM 024546 Homo sapiens hypothetical protein FLJ13449 (FLJ13449), mRNA NM 024534 Homo sapiens hypothetical protein FLJ12684 (FLJ12684), mRNA NM 024534 Homo sapiens hypothetical protein FLJ12584 (FLJ12684), mRNA NM 024526 Homo sapiens hypothetical protein FLJ1252724 (FLJ21522), mRNA NM 024526 Homo sapiens hypothetical protein FLJ21522 (FLJ21522), mRNA NM 024521 Homo sapiens hypothetical protein FLJ21523 (FLJ2035), mRNA NM 024522 Homo sapiens hypothetical protein FLJ12650 (FLJ12650), mRNA NM 024516 Homo sapiens hypothetical protein MGC4666 (MGC4666), mRNA NM 024511 Homo sapiens hypothetical protein MGC4666 (MGC4666), mRNA NM 024507 Homo sapiens hypothetical protein MGC4066 (MGC4663), mRNA NM 024507 Homo sapiens hypothetical protein MGC10791 (MGC10791), mRNA NM 024345 Homo sapiens Phosphatidylglycerophosphate Synthase (PGS1), mRNA NM 024340 Homo sapiens hypothetical protein MGC10765 (MGC10765), mRNA NM 024340 Homo sapiens hypothetical protein MGC479 (MGC4179), mRNA NM 024320 Homo sapiens hypothetical protein MGC4365 (MGC4365), mRNA NM 024321 Homo sapiens hypothetical protein MGC4179 (MGC11279), mRNA NM 024321 Homo sapiens hypothetical protein MGC4170 (MGC4170), mRNA NM 024321 Homo sapiens hypothetical protein MGC4170 (MGC4170), mRNA NM 024308 Homo sapiens hypothetical protein MGC4170 (MGC4170), mRNA NM 024309 Homo sapiens hypothetical protein MGC4170 (MGC4170), mRNA NM 024301 Homo sapiens hypothetical protein MGC4170 (MGC4170), mRNA NM 024301 Homo sapiens hypothetical protein MGC4170 (MGC4170), mRNA NM 024301 Homo sapiens hypothetical protein MGC4170 (MGC4171), mRNA NM 02401 Homo sapiens hypothetical protein MGC4707 (MGC4707), mRNA NM 02409 Homo sapiens hypothetical protein MGC3067 (MGC3067), mRNA NM 02409 Homo sapiens hypothetical protein MGC3067 (MGC3077), mRNA NM 02409 Homo sapiens hypothetical protein MGC		Homo sapiens hypothetical protein FLJ13119 (FLJ13119), mRNA
NM 024556 Homo sapiens hypothetical protein FLJ21103 (FLJ21103), mRNA NM 024552 Homo sapiens hypothetical protein FLJ12089 (FLJ12089), mRNA NM 024546 Homo sapiens hypothetical protein FLJ13449 (FLJ13449), mRNA NM 024534 Homo sapiens hypothetical protein FLJ12084 (FLJ12684), mRNA NM 024532 Homo sapiens hypothetical protein FLJ212724 (FLJ22724), mRNA NM 024526 Homo sapiens hypothetical protein FLJ21522 (FLJ21522), mRNA NM 024526 Homo sapiens hypothetical protein FLJ21522 (FLJ21522), mRNA NM 024526 Homo sapiens hypothetical protein FLJ21522 (FLJ21522), mRNA NM 024521 Homo sapiens hypothetical protein FLJ212035 (FLJ22635), mRNA NM 024516 Homo sapiens hypothetical protein MGC4606 (MGC4606), mRNA NM 024516 Homo sapiens hypothetical protein MGC4663 (MGC4663), mRNA NM 024517 Homo sapiens hypothetical protein MGC4663 (MGC4663), mRNA NM 015288 Homo sapiens KIAA0239 protein (KIAA0239), mRNA NM 024349 Homo sapiens Phosphatidylglycerophosphate Synthase (PGS1), mRNA NM 024340 Homo sapiens hypothetical protein MGC10765 (MGC10765), mRNA NM 024330 Homo sapiens hypothetical protein MGC4179 (MGC4179), mRNA NM 024321 Homo sapiens hypothetical protein MGC4179 (MGC4179), mRNA NM 024321 Homo sapiens hypothetical protein MGC11279 (MGC11279), mRNA NM 024321 Homo sapiens hypothetical protein MGC11279 (MGC11279), mRNA NM 024321 Homo sapiens hypothetical protein MGC4170 (MGC4170), mRNA NM 024308 Homo sapiens hypothetical protein MGC4170 (MGC4171), mRNA NM 024307 Homo sapiens hypothetical protein MGC4170 (MGC4171), mRNA NM 024308 Homo sapiens hypothetical protein MGC4170 (MGC4171), mRNA NM 024301 Homo sapiens hypothetical protein MGC4170 (MGC4171), mRNA NM 024301 Homo sapiens hypothetical protein MGC4170 (MGC4171), mRNA NM 02409 Homo sapiens hypothetical protein MGC40707 (MGC40707), mRNA NM 02409 Homo sapiens hypothetical protein MGC3067 (MGC3067), mRNA NM 02409 Homo sapiens hypothetical protein MGC3066 (MGC3069), mRNA NM 02409 Homo sapiens hypothetical protein MGC3066 (MGC3069), mRNA NM 024091 Homo sapiens hypothetical protein MGC3067 (MG		
NM 024546 NM 024546 NM 024546 NM 024546 NM 024534 NM 024534 NM 024534 NM 024532 NM 024526 NM 024526 NM 024526 NM 024526 NM 024528 NM 024528 NM 024528 NM 024528 NM 024528 NM 024529 NM 024520 NM 024520 NM 024521 NM 024521 NM 024521 NM 024522 NM 024522 NM NM 024522 NM 024522 NM 024522 NM 024524 NM 024524 NM 024526 NM 024526 NM 024526 NM 024527 NM 024527 NM 024528 NM 024528 NM 024528 NM 024529 NM 024520 NM 024521 NM 024516 NM 024516 NM 024516 NM 024517 NM 024517 NM 024517 NM 024518 NM 024518 NM 024518 NM 024519 NM 024519 NM 024510 NM 024511 NM 024511 NM 024512 NM 024512 NM 024513 NM 024514 NM 024514 NM 024515 NM 024515 NM 024516 NM 024516 NM 024516 NM 024517 NM 024517 NM 024518 NM 024518 NM 024519 NM 024510 NM 024510 NM 024510 NM 024510 NM 024510 NM 024310 NM 024310 NM 024321 NM 024320 NM 024320 NM 024321 NM 024322 NM 024323 NM 024321 NM 024324 NM 024325 NM 024326 NM 024327 NM 024327 NM 024328 NM 024328 NM 024329 NM 024321 NM 024329 NM 024329 NM 024329 NM 024321 NM 024329 NM 024321 NM 024325 NM 024327 NM 024328 NM 024328 NM 024329 NM 024328 NM 024329 NM 024329 NM 024329 NM 024329 NM 024329 NM 024329 NM 024328 NM 024434 NM 024436 NM 024436 NM 024436 NM 024437 NM 024437 NM 024438 NM 024448 NM 024467 NM 024092 NM 024093 NM 024094 NM 024094 NM 024094 NM		
NM 024524 Homo sapiens hypothetical protein FLJ13449 (FLJ13449), mRNA NM 024532 Homo sapiens hypothetical protein FLJ12684 (FLJ12684), mRNA NM 024522 Homo sapiens hypothetical protein FLJ22724 (FLJ2724), mRNA NM 024523 Homo sapiens hypothetical protein FLJ21522 (FLJ21522), mRNA NM 024523 Homo sapiens hypothetical protein FLJ21522 (FLJ21522), mRNA NM 024522 Homo sapiens hypothetical protein FLJ21650 (FLJ12650), mRNA NM 024522 Homo sapiens hypothetical protein MGC4606 (MGC4606), mRNA NM 024514 Homo sapiens hypothetical protein MGC4603 (MGC4663), mRNA NM 024514 Homo sapiens hypothetical protein MGC4603 (MGC4663), mRNA NM 024519 Homo sapiens hypothetical protein MGC10791 (MGC10791), mRNA NM 015288 Homo sapiens KIAA0239 protein (KIAA0239), mRNA NM 024419 Homo sapiens Phosphatidylglycerophosphate Synthase (PGS1), mRNA NM 024345 Homo sapiens hypothetical protein MGC10765 (MGC10765), mRNA NM 024340 Homo sapiens hypothetical protein MGC4179 (MGC4179), mRNA NM 024330 Homo sapiens hypothetical protein MGC10765 (MGC10765), mRNA NM 024326 Homo sapiens hypothetical protein MGC11279 (MGC11279), mRNA NM 024312 Homo sapiens hypothetical protein MGC10433 (MGC10433), mRNA NM 024312 Homo sapiens hypothetical protein MGC1470 (MGC4170), mRNA NM 024308 Homo sapiens hypothetical protein MGC1470 (MGC4170), mRNA NM 024307 Homo sapiens hypothetical protein MGC4171 (MGC4171), mRNA NM 024308 Homo sapiens hypothetical protein MGC4171 (MGC4171), mRNA NM 024307 Homo sapiens hypothetical protein MGC3067 (MGC3067), mRNA NM 024307 Homo sapiens hypothetical protein MGC4271 (MGC4171), mRNA NM 024308 Homo sapiens hypothetical protein MGC477 (MGC477), mRNA NM 024090 Homo sapiens hypothetical protein MGC3067 (MGC3067), mRNA NM 024091 Homo sapiens hypothetical protein MGC3067 (MGC3067), mRNA NM 024092 Homo sapiens hypothetical protein MGC4277 (MGC4827), mRNA NM 024094 Homo sapiens hypothetical protein MGC4277 (MGC4827), mRNA NM 024094 Homo sapiens hypothetical protein MGC3196 (MGC3306), mRNA NM 024097 Homo sapiens hypothetical protein MGC3196 (MGC3306		
NM 024524 Homo sapiens hypothetical protein FLJ12684 (FLJ12684), mRNA NM 024526 Homo sapiens hypothetical protein FLJ22724 (FLJ22724), mRNA NM 024526 Homo sapiens hypothetical protein FLJ21522 (FLJ21522), mRNA NM 024523 Homo sapiens hypothetical protein FLJ21522 (FLJ21522), mRNA NM 024524 Homo sapiens hypothetical protein FLJ21650 (FLJ12650), mRNA NM 024516 Homo sapiens hypothetical protein MGC4606 (MGC4606), mRNA NM 024514 Homo sapiens hypothetical protein MGC4606 (MGC4663), mRNA NM 024507 Homo sapiens hypothetical protein MGC10791 (MGC10791), mRNA NM 015288 Homo sapiens KIAA0239 protein (KIAA0239), mRNA NM 024419 Homo sapiens Phosphatidylglycerophosphate Synthase (PGS1), mRNA NM 024345 Homo sapiens hypothetical protein MGC10765 (MGC10765), mRNA NM 024340 Homo sapiens hypothetical protein MGC4179 (MGC4179), mRNA NM 024330 Homo sapiens hypothetical protein MGC4179 (MGC4179), mRNA NM 024321 Homo sapiens hypothetical protein MGC4365 (MGC4365), mRNA NM 024312 Homo sapiens hypothetical protein MGC107433 (MGC10433), mRNA NM 024312 Homo sapiens hypothetical protein MGC4170 (MGC4170), mRNA NM 024307 Homo sapiens hypothetical protein MGC4170 (MGC4171), mRNA NM 024307 Homo sapiens hypothetical protein MGC4171 (MGC4171), mRNA NM 024307 Homo sapiens hypothetical protein MGC4171 (MGC4171), mRNA NM 024307 Homo sapiens hypothetical protein MGC4171 (MGC4171), mRNA NM 024301 Homo sapiens hypothetical protein MGC4170 (MGC4171), mRNA NM 024301 Homo sapiens hypothetical protein MGC4170 (MGC4171), mRNA NM 024301 Homo sapiens hypothetical protein MGC4170 (MGC4171), mRNA NM 024301 Homo sapiens hypothetical protein MGC4170 (MGC4707), mRNA NM 024304 Homo sapiens hypothetical protein MGC4171 (MGC4707), mRNA NM 024095 Homo sapiens hypothetical protein MGC3067 (MGC3067), mRNA NM 024094 Homo sapiens hypothetical protein MGC5306 (MGC306), mRNA NM 024099 Homo sapiens hypothetical protein MGC2707 (MGC4707), mRNA NM 024091 Homo sapiens hypothetical protein MGC2306 (MGC306), mRNA NM 024092 Homo sapiens hypothetical protein MGC2718 (MGC3196), m		Homo sapiens hypothetical protein FLJ12089 (FLJ12089), mRNA
NM 024522 Homo sapiens hypothetical protein FLJ22724 (FLJ22724), mRNA NM 024523 Homo sapiens hypothetical protein FLJ21522 (FLJ21522), mRNA NM 024524 Homo sapiens hypothetical protein FLJ212035 (FLJ22035), mRNA NM 024516 Homo sapiens hypothetical protein MGC4606 (MGC4606), mRNA NM 024516 Homo sapiens hypothetical protein MGC4663 (MGC4663), mRNA NM 024517 Homo sapiens hypothetical protein MGC4663 (MGC4663), mRNA NM 024518 Homo sapiens hypothetical protein MGC10791 (MGC10791), mRNA NM 015288 Homo sapiens Phosphatidylglycerophosphate Synthase (PGS1), mRNA NM 024349 Homo sapiens hypothetical protein MGC10765 (MGC10765), mRNA NM 024340 Homo sapiens hypothetical protein MGC4179 (MGC4179), mRNA NM 024340 Homo sapiens hypothetical protein MGC4179 (MGC4179), mRNA NM 024320 Homo sapiens hypothetical protein MGC41279 (MGC4179), mRNA NM 024321 Homo sapiens hypothetical protein MGC10433 (MGC10433), mRNA NM 024312 Homo sapiens hypothetical protein MGC4170 (MGC4170), mRNA NM 024308 Homo sapiens hypothetical protein MGC4172 (MGC4172), mRNA NM 024307 Homo sapiens hypothetical protein MGC4171 (MGC4171), mRNA NM 024308 Homo sapiens hypothetical protein MGC4171 (MGC4171), mRNA NM 024309 Homo sapiens hypothetical protein MGC4170 (MGC4171), mRNA NM 024116 Homo sapiens hypothetical protein MGC4170 (MGC4171), mRNA NM 024116 Homo sapiens hypothetical protein MGC4707 (MGC4171), mRNA NM 024116 Homo sapiens hypothetical protein MGC4707 (MGC4171), mRNA NM 024116 Homo sapiens hypothetical protein MGC4707 (MGC4077), mRNA NM 024114 Homo sapiens hypothetical protein MGC53067 (MGC3067), mRNA NM 024099 Homo sapiens hypothetical protein MGC4707 (MGC4707), mRNA NM 024091 Homo sapiens hypothetical protein MGC3196 (MGC306), mRNA NM 024092 Homo sapiens hypothetical protein MGC3196 (MGC306), mRNA NM 024094 Homo sapiens hypothetical protein MGC3196 (MGC3196), mRNA NM 024094 Homo sapiens hypothetical protein MGC3196 (MGC3196), mRNA NM 024064 Homo sapiens hypothetical protein MGC3196 (MGC33547), mRNA NM 024063 Homo sapiens hypothetical protein MGC3196 (M		Homo sapiens hypothetical protein FLJ13449 (FLJ13449), mRNA
NM 024526 Homo sapiens hypothetical protein FLJ21522 (FLJ21522), mRNA NM 024523 Homo sapiens hypothetical protein FLJ2035 (FLJ22035), mRNA NM 024522 Homo sapiens hypothetical protein FLJ12650 (FLJ12650), mRNA NM 024516 Homo sapiens hypothetical protein MGC4606 (MGC4606), mRNA NM 024514 Homo sapiens hypothetical protein MGC4663 (MGC4663), mRNA NM 024507 Homo sapiens hypothetical protein MGC10791 (MGC10791), mRNA NM 015288 Homo sapiens KIAA0239 protein (KIAA0239), mRNA NM 024419 Homo sapiens Phosphatidylglycerophosphate Synthase (PGS1), mRNA NM 024345 Homo sapiens hypothetical protein MGC10765 (MGC10765), mRNA NM 024340 Homo sapiens hypothetical protein MGC4179 (MGC4179), mRNA NM 024326 Homo sapiens hypothetical protein MGC4179 (MGC4179), mRNA NM 024321 Homo sapiens hypothetical protein MGC10433 (MGC10433), mRNA NM 024321 Homo sapiens hypothetical protein MGC11279 (MGC11279), mRNA NM 024308 Homo sapiens hypothetical protein MGC4170 (MGC4170), mRNA NM 024309 Homo sapiens hypothetical protein MGC4172 (MGC4172), mRNA NM 024309 Homo sapiens hypothetical protein MGC4171 (MGC4171), mRNA NM 024091 Homo sapiens hypothetical protein MGC4171 (MGC4171), mRNA NM 024092 Homo sapiens hypothetical protein MGC4171 (MGC4171), mRNA NM 024114 Homo sapiens hypothetical protein MGC3067 (MGC3067), mRNA NM 024114 Homo sapiens SLC2A4 regulator (SLC2A4RG), mRNA NM 024114 Homo sapiens hypothetical protein MGC5306 (MGC5306), mRNA NM 024114 Homo sapiens hypothetical protein MGC4827 (MGC4827), mRNA NM 024114 Homo sapiens hypothetical protein MGC5306 (MGC5306), mRNA NM 024099 Homo sapiens hypothetical protein MGC5306 (MGC5508), mRNA NM 024094 Homo sapiens hypothetical protein MGC5306 (MGC5508), mRNA NM 024095 Homo sapiens hypothetical protein MGC3196 (MGC3196), mRNA NM 024096 Homo sapiens hypothetical protein MGC3196 (MGC3196), mRNA NM 024064 Homo sapiens hypothetical protein MGC3196 (MGC3196), mRNA NM 024064 Homo sapiens hypothetical protein MGC3196 (MGC335), mRNA NM 024064 Homo sapiens hypothetical protein MGC3196 (MGC33547), mRNA NM 024063		Homo sapiens hypothetical protein FLJ12684 (FLJ12684), mRNA
NM 024523 Homo sapiens hypothetical protein FLI22035 (FLI22035), mRNA NM 024512 Homo sapiens hypothetical protein FLJ12650 (FLJ12650), mRNA NM 024514 Homo sapiens hypothetical protein MGC4603 (MGC4606), mRNA NM 024514 Homo sapiens hypothetical protein MGC4603 (MGC4603), mRNA NM 024507 Homo sapiens hypothetical protein MGC10791 (MGC10791), mRNA NM 015288 Homo sapiens KIAA0239 protein (KIAA0239), mRNA NM 024419 Homo sapiens Phosphatidylglycerophosphate Synthase (PGS1), mRNA NM 024345 Homo sapiens hypothetical protein MGC10765 (MGC10765), mRNA NM 024340 Homo sapiens hypothetical protein MGC4179 (MGC4179), mRNA NM 024330 Homo sapiens hypothetical protein MGC4365 (MGC4365), mRNA NM 024326 Homo sapiens hypothetical protein MGC11279 (MGC11279), mRNA NM 024321 Homo sapiens hypothetical protein MGC10433 (MGC10433), mRNA NM 024312 Homo sapiens hypothetical protein MGC4170 (MGC4170), mRNA NM 024308 Homo sapiens hypothetical protein MGC4170 (MGC4170), mRNA NM 024307 Homo sapiens hypothetical protein MGC4171 (MGC4171), mRNA NM 024295 Homo sapiens hypothetical protein MGC3067 (MGC3067), mRNA NM 024092 Homo sapiens SLC2A4 regulator (SLC2A4RG), mRNA NM 024114 Homo sapiens hypothetical protein MGC3067 (MGC3067), mRNA NM 024113 Homo sapiens hypothetical protein MGC4707 (MGC4707), mRNA NM 024099 Homo sapiens hypothetical protein MGC4707 (MGC4707), mRNA NM 024099 Homo sapiens hypothetical protein MGC4707 (MGC4707), mRNA NM 024099 Homo sapiens hypothetical protein MGC4707 (MGC4707), mRNA NM 024091 Homo sapiens hypothetical protein MGC4707 (MGC4707), mRNA NM 024092 Homo sapiens hypothetical protein MGC3066 (MGC5306), mRNA NM 024094 Homo sapiens hypothetical protein MGC3067 (MGC3067), mRNA NM 024094 Homo sapiens hypothetical protein MGC3067 (MGC3067), mRNA NM 024094 Homo sapiens hypothetical protein MGC3196 (MGC3306), mRNA NM 024094 Homo sapiens hypothetical protein MGC3196 (MGC3305), mRNA NM 024064 Homo sapiens hypothetical protein MGC3196 (MGC3335), mRNA NM 024063 Homo sapiens hypothetical protein MGC3347 (MGC3347), mRNA NM 024063		
NM 024512 Homo sapiens hypothetical protein FLJ12650 (FLJ12650), mRNA NM 024516 Homo sapiens hypothetical protein MGC4606 (MGC4606), mRNA NM 024517 Homo sapiens hypothetical protein MGC10791 (MGC10791), mRNA NM 024507 Homo sapiens hypothetical protein MGC10791 (MGC10791), mRNA NM 015288 Homo sapiens RIAA0239 protein (KIAA0239), mRNA NM 024419 Homo sapiens Phosphatidylglycerophosphate Synthase (PGS1), mRNA NM 024345 Homo sapiens hypothetical protein MGC10765 (MGC10765), mRNA NM 024340 Homo sapiens hypothetical protein MGC4179 (MGC4179), mRNA NM 024330 Homo sapiens hypothetical protein MGC4365 (MGC4365), mRNA NM 024326 Homo sapiens hypothetical protein MGC11279 (MGC11279), mRNA NM 024321 Homo sapiens hypothetical protein MGC10433 (MGC10433), mRNA NM 024312 Homo sapiens hypothetical protein MGC4170 (MGC4170), mRNA NM 024308 Homo sapiens hypothetical protein MGC4172 (MGC4172), mRNA NM 024307 Homo sapiens hypothetical protein MGC4171 (MGC4171), mRNA NM 024295 Homo sapiens hypothetical protein MGC3067 (MGC3067), mRNA NM 024062 Homo sapiens SLC2A4 regulator (SLC2A4RG), mRNA NM 018491 Homo sapiens SLC2A4 regulator (SLC2A4RG), mRNA NM 024114 Homo sapiens hypothetical protein MGC5306 (MGC5306), mRNA NM 024114 Homo sapiens hypothetical protein MGC4277 (MGC4827), mRNA NM 024099 Homo sapiens hypothetical protein MGC4477 (MGC4707), mRNA NM 024099 Homo sapiens hypothetical protein MGC4277 (MGC4707), mRNA NM 024099 Homo sapiens hypothetical protein MGC4277 (MGC477), mRNA NM 024094 Homo sapiens hypothetical protein MGC3196 (MGC3365), mRNA NM 024094 Homo sapiens hypothetical protein MGC2118 (MGC2178), mRNA NM 024067 Homo sapiens hypothetical protein MGC2367 (MGC2335), mRNA NM 024063 Homo sapiens hypothetical protein MGC23071 (MGC2718), mRNA		
NM_024516 Homo sapiens hypothetical protein MGC4606 (MGC4606), mRNA NM_024514 Homo sapiens hypothetical protein MGC4603 (MGC4603), mRNA NM_024507 Homo sapiens hypothetical protein MGC10791 (MGC10791), mRNA NM_015288 Homo sapiens KIAA0239 protein (KIAA0239), mRNA NM_024419 Homo sapiens Phosphatidylglycerophosphate Synthase (PGS1), mRNA NM_024341 Homo sapiens hypothetical protein MGC10765 (MGC10765), mRNA NM_024344 Homo sapiens hypothetical protein MGC4179 (MGC4179), mRNA NM_024330 Homo sapiens hypothetical protein MGC4179 (MGC4179), mRNA NM_024326 Homo sapiens hypothetical protein MGC10433 (MGC10433), mRNA NM_024321 Homo sapiens hypothetical protein MGC10433 (MGC10433), mRNA NM_024312 Homo sapiens hypothetical protein MGC4170 (MGC4170), mRNA NM_024308 Homo sapiens hypothetical protein MGC4172 (MGC4172), mRNA NM_024307 Homo sapiens hypothetical protein MGC4171 (MGC4171), mRNA NM_024295 Homo sapiens hypothetical protein MGC3067 (MGC3067), mRNA NM_024094 Homo sapiens SLC2A4 regulator (SLC2A4RG), mRNA NM_024116 Homo sapiens hypothetical protein MGC5306 (MGC5306), mRNA NM_024114 Homo sapiens hypothetical protein MGC4827 (MGC4827), mRNA NM_024109 Homo sapiens hypothetical protein MGC4707 (MGC4707), mRNA NM_024099 Homo sapiens hypothetical protein MGC3067 (MGC4827), mRNA NM_024099 Homo sapiens hypothetical protein MGC3067 (MGC4827), mRNA NM_024099 Homo sapiens hypothetical protein MGC3067 (MGC4707), mRNA NM_024099 Homo sapiens hypothetical protein MGC3196 (MGC3196), mRNA NM_024091 Homo sapiens hypothetical protein MGC2365 (MGC5508), mRNA NM_024094 Homo sapiens hypothetical protein MGC2835 (MGC5508), mRNA NM_024064 Homo sapiens hypothetical protein MGC2835 (MGC2835), mRNA NM_024067 Homo sapiens hypothetical protein MGC2835 (MGC2835), mRNA NM_024063 Homo sapiens hypothetical protein MGC2836 (MGC5347), mRNA NM_024063 Homo sapiens hypothetical protein MGC3477 (MGC3477), mRNA		
NM 024514 Homo sapiens hypothetical protein MGC4663 (MGC4663), mRNA NM 024507 Homo sapiens KIAA0239 protein (KIAA0239), mRNA NM 015288 Homo sapiens KIAA0239 protein (KIAA0239), mRNA NM 024419 Homo sapiens Phosphatidylglycerophosphate Synthase (PGS1), mRNA NM 024345 Homo sapiens hypothetical protein MGC10765 (MGC10765), mRNA NM 024346 Homo sapiens hypothetical protein MGC4179 (MGC4179), mRNA NM 024340 Homo sapiens hypothetical protein MGC4365 (MGC4365), mRNA NM 024320 Homo sapiens hypothetical protein MGC11279 (MGC11279), mRNA NM 024321 Homo sapiens hypothetical protein MGC10433 (MGC10433), mRNA NM 024312 Homo sapiens hypothetical protein MGC4170 (MGC4170), mRNA NM 024308 Homo sapiens hypothetical protein MGC4172 (MGC4172), mRNA NM 024307 Homo sapiens hypothetical protein MGC4171 (MGC4171), mRNA NM 024295 Homo sapiens hypothetical protein MGC3067 (MGC3067), mRNA NM 020062 Homo sapiens SLC2A4 regulator (SLC2A4R6), mRNA NM 024116 Homo sapiens hypothetical protein MGC5306 (MGC5306), mRNA NM 024114 Homo sapiens hypothetical protein MGC4827 (MGC4827), mRNA NM 024113 Homo sapiens hypothetical protein MGC4707 (MGC4707), mRNA NM 024099 Homo sapiens hypothetical protein MGC4707 (MGC4707), mRNA NM 024092 Homo sapiens hypothetical protein MGC5308 (MGC5308), mRNA NM 024094 Homo sapiens hypothetical protein MGC4707 (MGC4707), mRNA NM 024094 Homo sapiens hypothetical protein MGC2477 (MGC2477), mRNA NM 024094 Homo sapiens hypothetical protein MGC3196 (MGC3395), mRNA NM 024094 Homo sapiens hypothetical protein MGC3196 (MGC335), mRNA NM 024094 Homo sapiens hypothetical protein MGC3196 (MGC3505), mRNA NM 024094 Homo sapiens hypothetical protein MGC3196 (MGC335), mRNA NM 024064 Homo sapiens hypothetical protein MGC3198 (MGC3196), mRNA NM 024063 Homo sapiens hypothetical protein MGC3196 (MGC335), mRNA NM 024064 Homo sapiens hypothetical protein MGC3198 (MGC3196), mRNA NM 024064 Homo sapiens hypothetical protein MGC3198 (MGC3196), mRNA		
NM_024507 Homo sapiens hypothetical protein MGC10791 (MGC10791), mRNA NM_015288 Homo sapiens KIAA0239 protein (KIAA0239), mRNA NM_024419 Homo sapiens Phosphatidylglycerophosphate Synthase (PGS1), mRNA NM_024345 Homo sapiens hypothetical protein MGC10765 (MGC10765), mRNA NM_024340 Homo sapiens hypothetical protein MGC4179 (MGC4179), mRNA NM_024330 Homo sapiens hypothetical protein MGC4365 (MGC4365), mRNA NM_024326 Homo sapiens hypothetical protein MGC10433 (MGC10433), mRNA NM_024321 Homo sapiens hypothetical protein MGC10433 (MGC10433), mRNA NM_024312 Homo sapiens hypothetical protein MGC4170 (MGC4170), mRNA NM_024308 Homo sapiens hypothetical protein MGC4172 (MGC4172), mRNA NM_024307 Homo sapiens hypothetical protein MGC4171 (MGC4171), mRNA NM_024295 Homo sapiens hypothetical protein MGC3067 (MGC3067), mRNA NM_024062 Homo sapiens SLC2A4 regulator (SLC2A4RG), mRNA NM_018491 Homo sapiens COBW-like protein (LOC55871), mRNA NM_024116 Homo sapiens hypothetical protein MGC306 (MGC5306), mRNA NM_024111 Homo sapiens hypothetical protein MGC4827 (MGC4827), mRNA NM_024113 Homo sapiens hypothetical protein MGC4707 (MGC4707), mRNA NM_024099 Homo sapiens hypothetical protein MGC4707 (MGC4707), mRNA NM_024099 Homo sapiens hypothetical protein MGC5508 (MGC5508), mRNA NM_024092 Homo sapiens hypothetical protein MGC5508 (MGC5508), mRNA NM_024094 Homo sapiens hypothetical protein MGC5306 (MGC5306), mRNA NM_024094 Homo sapiens hypothetical protein MGC5508 (MGC5508), mRNA NM_024094 Homo sapiens hypothetical protein MGC5508 (MGC5508), mRNA NM_024096 Homo sapiens hypothetical protein MGC5306 (MGC3196), mRNA NM_024097 Homo sapiens hypothetical protein MGC2477 (MGC2477), mRNA NM_024067 Homo sapiens hypothetical protein MGC3196 (MGC3196), mRNA NM_024063 Homo sapiens hypothetical protein MGC2718 (MGC2315), mRNA NM_024063 Homo sapiens hypothetical protein MGC5547 (MGC5347), mRNA		Homo sapiens hypothetical protein MGC4606 (MGC4606), mRNA
NM 015288 Homo sapiens KIAA0239 protein (KIAA0239), mRNA NM 024419 Homo sapiens Phosphatidylglycerophosphate Synthase (PGS1), mRNA NM 024345 Homo sapiens hypothetical protein MGC10765 (MGC10765), mRNA NM 024340 Homo sapiens hypothetical protein MGC4179 (MGC4179), mRNA NM 024330 Homo sapiens hypothetical protein MGC4365 (MGC4365), mRNA NM 024326 Homo sapiens hypothetical protein MGC11279 (MGC11279), mRNA NM 024321 Homo sapiens hypothetical protein MGC10433 (MGC10433), mRNA NM 024312 Homo sapiens hypothetical protein MGC4170 (MGC4170), mRNA NM 024308 Homo sapiens hypothetical protein MGC4172 (MGC4172), mRNA NM 024307 Homo sapiens hypothetical protein MGC4171 (MGC4171), mRNA NM 024295 Homo sapiens hypothetical protein MGC3067 (MGC3067), mRNA NM 020062 Homo sapiens SLC2A4 regulator (SLC2A4RG), mRNA NM 024116 Homo sapiens hypothetical protein MGC5306 (MGC5306), mRNA NM 024114 Homo sapiens hypothetical protein MGC4827 (MGC4827), mRNA NM 024113 Homo sapiens hypothetical protein MGC4707 (MGC4707), mRNA NM 024099 Homo sapiens hypothetical protein MGC2477 (MGC2477), mRNA NM 024099 Homo sapiens hypothetical protein MGC5508 (MGC5508), mRNA NM 024084 Homo sapiens hypothetical protein MGC2536 (MGC5508), mRNA NM 024084 Homo sapiens hypothetical protein MGC2538 (MGC5508), mRNA NM 024084 Homo sapiens hypothetical protein MGC23196 (MGC5306), mRNA NM 024084 Homo sapiens hypothetical protein MGC2718 (MGC2835), mRNA NM 024067 Homo sapiens hypothetical protein MGC2718 (MGC2718), mRNA NM 024063 Homo sapiens hypothetical protein MGC2718 (MGC2718), mRNA NM 024063 Homo sapiens hypothetical protein MGC2718 (MGC5547), mRNA		
NM 024419 Homo sapiens Phosphatidylglycerophosphate Synthase (PGS1), mRNA NM 024345 Homo sapiens hypothetical protein MGC10765 (MGC10765), mRNA NM 024340 Homo sapiens hypothetical protein MGC4179 (MGC4179), mRNA NM 024330 Homo sapiens hypothetical protein MGC4365 (MGC4365), mRNA NM 024326 Homo sapiens hypothetical protein MGC11279 (MGC11279), mRNA NM 024321 Homo sapiens hypothetical protein MGC10433 (MGC10433), mRNA NM 024312 Homo sapiens hypothetical protein MGC4170 (MGC4170), mRNA NM 024308 Homo sapiens hypothetical protein MGC4172 (MGC4172), mRNA NM 024307 Homo sapiens hypothetical protein MGC4171 (MGC4171), mRNA NM 024295 Homo sapiens hypothetical protein MGC3067 (MGC3067), mRNA NM 024062 Homo sapiens SLC2A4 regulator (SLC2A4RG), mRNA NM 024116 Homo sapiens hypothetical protein (LOC55871), mRNA NM 024114 Homo sapiens hypothetical protein MGC306 (MGC5306), mRNA NM 024114 Homo sapiens hypothetical protein MGC4827 (MGC4827), mRNA NM 024113 Homo sapiens hypothetical protein MGC4707 (MGC4707), mRNA NM 024099 Homo sapiens hypothetical protein MGC4707 (MGC4707), mRNA NM 024094 Homo sapiens hypothetical protein MGC4707 (MGC4707), mRNA NM 024084 Homo sapiens hypothetical protein MGC5508 (MGC5508), mRNA NM 024084 Homo sapiens hypothetical protein MGC3196 (MGC3196), mRNA NM 024067 Homo sapiens hypothetical protein MGC2835 (MGC2835), mRNA NM 024063 Homo sapiens hypothetical protein MGC2718 (MGC2718), mRNA NM 024063 Homo sapiens hypothetical protein MGC2718 (MGC2718), mRNA		Homo sapiens hypothetical protein MGC10791 (MGC10791), mRNA
NM 024345 Homo sapiens hypothetical protein MGC10765 (MGC10765), mRNA NM 024340 Homo sapiens hypothetical protein MGC4179 (MGC4179), mRNA NM 024330 Homo sapiens hypothetical protein MGC4365 (MGC4365), mRNA NM 024326 Homo sapiens hypothetical protein MGC11279 (MGC11279), mRNA NM 024321 Homo sapiens hypothetical protein MGC10433 (MGC10433), mRNA NM 024312 Homo sapiens hypothetical protein MGC4170 (MGC4170), mRNA NM 024308 Homo sapiens hypothetical protein MGC4172 (MGC4172), mRNA NM 024307 Homo sapiens hypothetical protein MGC4171 (MGC4171), mRNA NM 024295 Homo sapiens hypothetical protein MGC3067 (MGC3067), mRNA NM 024062 Homo sapiens SLC2A4 regulator (SLC2A4RG), mRNA NM 018491 Homo sapiens COBW-like protein (LOC55871), mRNA NM 024116 Homo sapiens hypothetical protein MGC306 (MGC5306), mRNA NM 024113 Homo sapiens hypothetical protein MGC4827 (MGC4827), mRNA NM 024099 Homo sapiens hypothetical protein MGC4707 (MGC4707), mRNA NM 024092 Homo sapiens hypothetical protein MGC2477 (MGC4770), mRNA NM 024092 Homo sapiens hypothetical protein MGC5508 (MGC5508), mRNA NM 024084 Homo sapiens hypothetical protein MGC3196 (MGC53196), mRNA NM 024072 Homo sapiens hypothetical protein MGC2835 (MGC2835), mRNA NM 024067 Homo sapiens hypothetical protein MGC2718 (MGC2718), mRNA NM 024063 Homo sapiens hypothetical protein MGC2718 (MGC2718), mRNA NM 024063 Homo sapiens hypothetical protein MGC5347 (MGC5347), mRNA		
NM 024340 Homo sapiens hypothetical protein MGC4179 (MGC4179), mRNA NM 024330 Homo sapiens hypothetical protein MGC4365 (MGC4365), mRNA NM 024326 Homo sapiens hypothetical protein MGC11279 (MGC11279), mRNA NM 024321 Homo sapiens hypothetical protein MGC10433 (MGC10433), mRNA NM 024312 Homo sapiens hypothetical protein MGC4170 (MGC4170), mRNA NM 024308 Homo sapiens hypothetical protein MGC4172 (MGC4172), mRNA NM 024307 Homo sapiens hypothetical protein MGC4171 (MGC4171), mRNA NM 024295 Homo sapiens hypothetical protein MGC3067 (MGC3067), mRNA NM 020062 Homo sapiens SLC2A4 regulator (SLC2A4RG), mRNA NM 018491 Homo sapiens COBW-like protein (LOC55871), mRNA NM 024116 Homo sapiens hypothetical protein MGC4827 (MGC4827), mRNA NM 024113 Homo sapiens hypothetical protein MGC4707 (MGC4707), mRNA NM 024099 Homo sapiens hypothetical protein MGC4707 (MGC4707), mRNA NM 024092 Homo sapiens hypothetical protein MGC5508 (MGC5508), mRNA NM 024084 Homo sapiens hypothetical protein MGC5508 (MGC5508), mRNA NM 024072 Homo sapiens hypothetical protein MGC3196 (MGC3196), mRNA NM 024067 Homo sapiens hypothetical protein MGC2835 (MGC2835), mRNA NM 024063 Homo sapiens hypothetical protein MGC2718 (MGC2718), mRNA NM 024063 Homo sapiens hypothetical protein MGC5347 (MGC5347), mRNA NM 024063 Homo sapiens hypothetical protein MGC5347 (MGC5347), mRNA		Homo sapiens Phosphatidylglycerophosphate Synthase (PGS1), mRNA
NM 024330 Homo sapiens hypothetical protein MGC4365 (MGC4365), mRNA NM_024326 Homo sapiens hypothetical protein MGC11279 (MGC11279), mRNA NM_024321 Homo sapiens hypothetical protein MGC10433 (MGC10433), mRNA NM_024312 Homo sapiens hypothetical protein MGC4170 (MGC4170), mRNA NM_024308 Homo sapiens hypothetical protein MGC4172 (MGC4172), mRNA NM_024307 Homo sapiens hypothetical protein MGC4171 (MGC4171), mRNA NM_024295 Homo sapiens hypothetical protein MGC3067 (MGC3067), mRNA NM_020062 Homo sapiens SLC2A4 regulator (SLC2A4RG), mRNA NM_018491 Homo sapiens COBW-like protein (LOC55871), mRNA NM_024116 Homo sapiens hypothetical protein MGC5306 (MGC5306), mRNA NM_024114 Homo sapiens hypothetical protein MGC4827 (MGC4827), mRNA NM_024113 Homo sapiens hypothetical protein MGC4707 (MGC4707), mRNA NM_024099 Homo sapiens hypothetical protein MGC2477 (MGC2477), mRNA NM_024092 Homo sapiens hypothetical protein MGC5508 (MGC5508), mRNA NM_024084 Homo sapiens hypothetical protein MGC3196 (MGC3196), mRNA NM_024072 Homo sapiens hypothetical protein MGC2835 (MGC2835), mRNA NM_024067 Homo sapiens hypothetical protein MGC2718 (MGC2718), mRNA NM_024063 Homo sapiens hypothetical protein MGC5347 (MGC5347), mRNA NM_024063 Homo sapiens hypothetical protein MGC5347 (MGC5347), mRNA		
NM 024326 Homo sapiens hypothetical protein MGC11279 (MGC11279), mRNA NM 024321 Homo sapiens hypothetical protein MGC10433 (MGC10433), mRNA NM 024312 Homo sapiens hypothetical protein MGC4170 (MGC4170), mRNA NM 024308 Homo sapiens hypothetical protein MGC4172 (MGC4172), mRNA NM 024307 Homo sapiens hypothetical protein MGC4171 (MGC4171), mRNA NM 024295 Homo sapiens hypothetical protein MGC3067 (MGC3067), mRNA NM 020062 Homo sapiens SLC2A4 regulator (SLC2A4RG), mRNA NM 018491 Homo sapiens COBW-like protein (LOC55871), mRNA NM 024116 Homo sapiens hypothetical protein MGC5306 (MGC5306), mRNA NM 024114 Homo sapiens hypothetical protein MGC4827 (MGC4827), mRNA NM 024113 Homo sapiens hypothetical protein MGC4707 (MGC4707), mRNA NM 024099 Homo sapiens hypothetical protein MGC2477 (MGC2477), mRNA NM 024084 Homo sapiens hypothetical protein MGC5508 (MGC5508), mRNA NM 024084 Homo sapiens hypothetical protein MGC3196 (MGC3196), mRNA NM 024072 Homo sapiens hypothetical protein MGC2835 (MGC2835), mRNA NM 024067 Homo sapiens hypothetical protein MGC2718 (MGC2718), mRNA NM 024063 Homo sapiens hypothetical protein MGC2718 (MGC2718), mRNA NM 024063 Homo sapiens hypothetical protein MGC5347 (MGC5347), mRNA		
NM 024321 Homo sapiens hypothetical protein MGC10433 (MGC10433), mRNA NM 024312 Homo sapiens hypothetical protein MGC4170 (MGC4170), mRNA NM 024308 Homo sapiens hypothetical protein MGC4172 (MGC4172), mRNA NM 024307 Homo sapiens hypothetical protein MGC4171 (MGC4171), mRNA NM 024295 Homo sapiens hypothetical protein MGC3067 (MGC3067), mRNA NM 020062 Homo sapiens SLC2A4 regulator (SLC2A4RG), mRNA NM 018491 Homo sapiens COBW-like protein (LOC55871), mRNA NM 024116 Homo sapiens hypothetical protein MGC5306 (MGC5306), mRNA NM 024114 Homo sapiens hypothetical protein MGC4827 (MGC4827), mRNA NM 024113 Homo sapiens hypothetical protein MGC4707 (MGC4707), mRNA NM 024099 Homo sapiens hypothetical protein MGC2477 (MGC2477), mRNA NM 024092 Homo sapiens hypothetical protein MGC5508 (MGC5508), mRNA NM 024084 Homo sapiens hypothetical protein MGC3196 (MGC3196), mRNA NM 024072 Homo sapiens hypothetical protein MGC2835 (MGC2835), mRNA NM 024067 Homo sapiens hypothetical protein MGC2718 (MGC2718), mRNA NM 024063 Homo sapiens hypothetical protein MGC5347 (MGC5347), mRNA		
NM 024312 Homo sapiens hypothetical protein MGC4170 (MGC4170), mRNA NM 024308 Homo sapiens hypothetical protein MGC4172 (MGC4172), mRNA NM 024307 Homo sapiens hypothetical protein MGC4171 (MGC4171), mRNA NM 024295 Homo sapiens hypothetical protein MGC3067 (MGC3067), mRNA NM 020062 Homo sapiens SLC2A4 regulator (SLC2A4RG), mRNA NM 018491 Homo sapiens COBW-like protein (LOC55871), mRNA NM 024116 Homo sapiens hypothetical protein MGC5306 (MGC5306), mRNA NM 024114 Homo sapiens hypothetical protein MGC4827 (MGC4827), mRNA NM 024113 Homo sapiens hypothetical protein MGC4707 (MGC4707), mRNA NM 024099 Homo sapiens hypothetical protein MGC2477 (MGC2477), mRNA NM 024092 Homo sapiens hypothetical protein MGC5508 (MGC5508), mRNA NM 024084 Homo sapiens hypothetical protein MGC3196 (MGC3196), mRNA NM 024072 Homo sapiens hypothetical protein MGC2835 (MGC2835), mRNA NM 024067 Homo sapiens hypothetical protein MGC2718 (MGC2718), mRNA NM 024063 Homo sapiens hypothetical protein MGC2718 (MGC2718), mRNA NM 024063 Homo sapiens hypothetical protein MGC5347 (MGC5347), mRNA		
NM 024308 Homo sapiens hypothetical protein MGC4172 (MGC4172), mRNA NM 024307 Homo sapiens hypothetical protein MGC4171 (MGC4171), mRNA NM 024295 Homo sapiens hypothetical protein MGC3067 (MGC3067), mRNA NM 020062 Homo sapiens SLC2A4 regulator (SLC2A4RG), mRNA NM 018491 Homo sapiens COBW-like protein (LOC55871), mRNA NM 024116 Homo sapiens hypothetical protein MGC5306 (MGC5306), mRNA NM 024114 Homo sapiens hypothetical protein MGC4827 (MGC4827), mRNA NM 024113 Homo sapiens hypothetical protein MGC4707 (MGC4707), mRNA NM 024099 Homo sapiens hypothetical protein MGC2477 (MGC2477), mRNA NM 024092 Homo sapiens hypothetical protein MGC5508 (MGC5508), mRNA NM 024084 Homo sapiens hypothetical protein MGC3196 (MGC3196), mRNA NM 024072 Homo sapiens hypothetical protein MGC2835 (MGC2835), mRNA NM 024067 Homo sapiens hypothetical protein MGC2718 (MGC2718), mRNA NM 024063 Homo sapiens hypothetical protein MGC5347 (MGC5347), mRNA NM 024063 Homo sapiens hypothetical protein MGC5347 (MGC5347), mRNA		
NM 024307 Homo sapiens hypothetical protein MGC4171 (MGC4171), mRNA NM 024295 Homo sapiens hypothetical protein MGC3067 (MGC3067), mRNA NM 020062 Homo sapiens SLC2A4 regulator (SLC2A4RG), mRNA NM 018491 Homo sapiens COBW-like protein (LOC55871), mRNA NM 024116 Homo sapiens hypothetical protein MGC5306 (MGC5306), mRNA NM 024114 Homo sapiens hypothetical protein MGC4827 (MGC4827), mRNA NM 024113 Homo sapiens hypothetical protein MGC4707 (MGC4707), mRNA NM 024099 Homo sapiens hypothetical protein MGC2477 (MGC2477), mRNA NM 024092 Homo sapiens hypothetical protein MGC5508 (MGC5508), mRNA NM 024084 Homo sapiens hypothetical protein MGC3196 (MGC3196), mRNA NM 024072 Homo sapiens hypothetical protein MGC2835 (MGC2835), mRNA NM 024067 Homo sapiens hypothetical protein MGC2718 (MGC2718), mRNA NM 024063 Homo sapiens hypothetical protein MGC5347 (MGC5347), mRNA		Homo sapiens hypothetical protein MGC4170 (MGC4170), mRNA
NM 024295 Homo sapiens hypothetical protein MGC3067 (MGC3067), mRNA  NM 020062 Homo sapiens SLC2A4 regulator (SLC2A4RG), mRNA  NM 018491 Homo sapiens COBW-like protein (LOC55871), mRNA  NM 024116 Homo sapiens hypothetical protein MGC5306 (MGC5306), mRNA  NM 024114 Homo sapiens hypothetical protein MGC4827 (MGC4827), mRNA  NM 024113 Homo sapiens hypothetical protein MGC4707 (MGC4707), mRNA  NM 024099 Homo sapiens hypothetical protein MGC2477 (MGC2477), mRNA  NM 024092 Homo sapiens hypothetical protein MGC5508 (MGC5508), mRNA  NM 024084 Homo sapiens hypothetical protein MGC3196 (MGC3196), mRNA  NM 024072 Homo sapiens hypothetical protein MGC2835 (MGC2835), mRNA  NM 024067 Homo sapiens hypothetical protein MGC2718 (MGC2718), mRNA  NM 024063 Homo sapiens hypothetical protein MGC5347 (MGC5347), mRNA		Homo sapiens hypothetical protein MGC4172 (MGC4172), mRNA
NM 020062 Homo sapiens SLC2A4 regulator (SLC2A4RG), mRNA  NM 018491 Homo sapiens COBW-like protein (LOC55871), mRNA  NM 024116 Homo sapiens hypothetical protein MGC5306 (MGC5306), mRNA  NM 024114 Homo sapiens hypothetical protein MGC4827 (MGC4827), mRNA  NM 024113 Homo sapiens hypothetical protein MGC4707 (MGC4707), mRNA  NM 024099 Homo sapiens hypothetical protein MGC2477 (MGC2477), mRNA  NM 024092 Homo sapiens hypothetical protein MGC5508 (MGC5508), mRNA  NM 024084 Homo sapiens hypothetical protein MGC3196 (MGC3196), mRNA  NM 024072 Homo sapiens hypothetical protein MGC2835 (MGC2835), mRNA  NM 024067 Homo sapiens hypothetical protein MGC2718 (MGC2718), mRNA  NM 024063 Homo sapiens hypothetical protein MGC5347 (MGC5347), mRNA		
NM 018491 Homo sapiens COBW-like protein (LOC55871), mRNA NM 024116 Homo sapiens hypothetical protein MGC5306 (MGC5306), mRNA NM 024114 Homo sapiens hypothetical protein MGC4827 (MGC4827), mRNA NM 024113 Homo sapiens hypothetical protein MGC4707 (MGC4707), mRNA NM 024099 Homo sapiens hypothetical protein MGC2477 (MGC2477), mRNA NM 024092 Homo sapiens hypothetical protein MGC5508 (MGC5508), mRNA NM 024084 Homo sapiens hypothetical protein MGC3196 (MGC3196), mRNA NM 024072 Homo sapiens hypothetical protein MGC2835 (MGC2835), mRNA NM 024067 Homo sapiens hypothetical protein MGC2718 (MGC2718), mRNA NM 024063 Homo sapiens hypothetical protein MGC5347 (MGC5347), mRNA		
NM 024116 Homo sapiens hypothetical protein MGC5306 (MGC5306), mRNA NM 024114 Homo sapiens hypothetical protein MGC4827 (MGC4827), mRNA NM 024113 Homo sapiens hypothetical protein MGC4707 (MGC4707), mRNA NM 024099 Homo sapiens hypothetical protein MGC2477 (MGC2477), mRNA NM 024092 Homo sapiens hypothetical protein MGC5508 (MGC5508), mRNA NM 024084 Homo sapiens hypothetical protein MGC3196 (MGC3196), mRNA NM 024072 Homo sapiens hypothetical protein MGC2835 (MGC2835), mRNA NM 024067 Homo sapiens hypothetical protein MGC2718 (MGC2718), mRNA NM 024063 Homo sapiens hypothetical protein MGC5347 (MGC5347), mRNA		Homo sapiens SLC2A4 regulator (SLC2A4RG), mRNA
NM 024114 Homo sapiens hypothetical protein MGC4827 (MGC4827), mRNA  NM 024113 Homo sapiens hypothetical protein MGC4707 (MGC4707), mRNA  NM 024099 Homo sapiens hypothetical protein MGC2477 (MGC2477), mRNA  NM 024092 Homo sapiens hypothetical protein MGC5508 (MGC5508), mRNA  NM 024084 Homo sapiens hypothetical protein MGC3196 (MGC3196), mRNA  NM 024072 Homo sapiens hypothetical protein MGC2835 (MGC2835), mRNA  NM 024067 Homo sapiens hypothetical protein MGC2718 (MGC2718), mRNA  NM 024063 Homo sapiens hypothetical protein MGC5347 (MGC5347), mRNA		
NM 024099 Homo sapiens hypothetical protein MGC4707 (MGC4707), mRNA NM 024099 Homo sapiens hypothetical protein MGC2477 (MGC2477), mRNA NM 024092 Homo sapiens hypothetical protein MGC5508 (MGC5508), mRNA NM 024084 Homo sapiens hypothetical protein MGC3196 (MGC3196), mRNA NM 024072 Homo sapiens hypothetical protein MGC2835 (MGC2835), mRNA NM 024067 Homo sapiens hypothetical protein MGC2718 (MGC2718), mRNA NM 024063 Homo sapiens hypothetical protein MGC5347 (MGC5347), mRNA		
NM 024099 Homo sapiens hypothetical protein MGC2477 (MGC2477), mRNA NM 024092 Homo sapiens hypothetical protein MGC5508 (MGC5508), mRNA NM 024084 Homo sapiens hypothetical protein MGC3196 (MGC3196), mRNA NM 024072 Homo sapiens hypothetical protein MGC2835 (MGC2835), mRNA NM 024067 Homo sapiens hypothetical protein MGC2718 (MGC2718), mRNA NM 024063 Homo sapiens hypothetical protein MGC5347 (MGC5347), mRNA		
NM 024092 Homo sapiens hypothetical protein MGC5508 (MGC5508), mRNA  NM 024084 Homo sapiens hypothetical protein MGC3196 (MGC3196), mRNA  NM 024072 Homo sapiens hypothetical protein MGC2835 (MGC2835), mRNA  NM 024067 Homo sapiens hypothetical protein MGC2718 (MGC2718), mRNA  NM 024063 Homo sapiens hypothetical protein MGC5347 (MGC5347), mRNA		
NM_024084 Homo sapiens hypothetical protein MGC3196 (MGC3196), mRNA NM_024072 Homo sapiens hypothetical protein MGC2835 (MGC2835), mRNA NM_024067 Homo sapiens hypothetical protein MGC2718 (MGC2718), mRNA NM_024063 Homo sapiens hypothetical protein MGC5347 (MGC5347), mRNA		Homo sapiens hypothetical protein MGC2477 (MGC2477), mRNA
NM 024072 Homo sapiens hypothetical protein MGC2835 (MGC2835), mRNA NM 024067 Homo sapiens hypothetical protein MGC2718 (MGC2718), mRNA NM 024063 Homo sapiens hypothetical protein MGC5347 (MGC5347), mRNA		Homo sapiens hypothetical protein MGC5508 (MGC5508), mRNA
NM_024067 Homo sapiens hypothetical protein MGC2718 (MGC2718), mRNA NM_024063 Homo sapiens hypothetical protein MGC5347 (MGC5347), mRNA		Homo sapiens hypothetical protein MGC3196 (MGC3196), mRNA
NM_024067 Homo sapiens hypothetical protein MGC2718 (MGC2718), mRNA NM_024063 Homo sapiens hypothetical protein MGC5347 (MGC5347), mRNA		Homo sapiens hypothetical protein MGC2835 (MGC2835), mRNA
1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1		Homo sapiens hypothetical protein MGC2718 (MGC2718), mRNA
313.4 (00.40.40   1.77		
The state of the s	NM_024040	Homo sapiens hypothetical protein MGC2491 (MGC2491), mRNA
NM_024036 Homo sapiens hypothetical protein MGC3103 (MGC3103), mRNA		Homo sapiens hypothetical protein MGC3103 (MGC3103), mRNA
NM_015450 Homo sapiens protection of telomeres 1 (POT1), mRNA		
NM_021249 Homo sapiens sorting nexin 6 (SNX6), mRNA		
NM_023932 Homo sapiens hypothetical protein MGC2487 (MGC2487), mRNA		Homo sapiens hypothetical protein MGC2487 (MGC2487), mRNA
NM_023930 Homo sapiens hypothetical protein MGC2376 (MGC2376), mRNA	NM_023930	Homo sapiens hypothetical protein MGC2376 (MGC2376), mRNA

	OVEZDS64C1040) mPNA
NM_014045	Homo sapiens DKFZP564C1940 protein (DKFZP564C1940), mRNA
NM_015533	Homo sapiens DKFZP586B1621 protein (DKFZP586B1621), mRNA
NM_023927	Homo sapiens hypothetical protein FLJ21313 (FLJ21313), mRNA
NM_023923	Homo sapiens hypothetical protein FLJ13171 (FLJ13171), mRNA
NM 019054	Homo sapiens hypothetical protein MGC5560 (MGC5560), mRNA
NIM 023070	Homo saniens hypothetical protein (LOC65243), mRNA
NM 023015	Homo conjens hypothetical protein FLJ21919 (FLJ21919), mRNA
NM_022899	Homo sapiens likely ortholog of mouse actin-related protein 8 homolog (S.
-	acception (FI I12034) mRNA
NM 022836	Homo sapiens DNA cross-link repair 1B (PSO2 homolog, S. cerevisiae)
- ]	(DCI PE1R) mRNA
NM 022831	Homo sapiens hypothetical protein FLJ12806 (FLJ12806), mRNA
NM 022828	Homo saniens hypothetical protein FLJ21940 (FLJ21940), mknA
NM_022822	Homo sapiens hypothetical protein FLJ12387 similar to kinesin light chain
112.2	(FI 112387) mRNA
NM 022784	Homo sapiens hypothetical protein FLJ12476 (FLJ12476), mRNA
NM 022783	Homo saniens hypothetical protein FLJ12428 (FLJ12428), mRNA
NM 022774	Homo saniens hypothetical protein FLJ21144 (FLJ21144), mRNA
NM 022765	Homo saniens hypothetical protein FLJ11937 (FLJ11937), mRNA
NM 022764	Homo saniens hypothetical protein FLJ12998 (FLJ12998), mRNA
NM 022758	Homo sapiens hypothetical protein FLJ22195 (FLJ22195), mRNA
NM 022753	Homo sapiens hypothetical protein FLJ12903 (FLJ12903), mRNA
NM 022749	Homo sapiens retinoic acid induced 16 (RAI16), mRNA
NM 022746	Homo sapiens hypothetical protein FLJ22390 (FLJ22390), mRNA
NM 022728	Homo sapiens neurogenic differentiation 6 (NEUROD6), mRNA
NM 022496	Homo saniens hypothetical protein FLJ13433 (FLJ13433), mRNA
NM 022490	Homo sapiens hypothetical protein FLJ13390 similar to PAF53 (FLJ13390),
NIVI_022490	mP N A
NM 022484	Homo saniens hypothetical protein FLJ13576 (FLJ13576), mRNA
NM 022483	Homo sapiens hypothetical protein FLJ21657 (FLJ21657), mRNA
NM 022473	Homo sapiens zinc finger protein 106 (ZFP106), mRNA
NM_022473	Homo sapiens hypothetical protein FLJ13057 similar to germ cell-less
NM_0224/1	(FLJ13057), mRNA
ND 4 022463	Homo saniens nubleoredovin 1 (NXN), mRNA
NM_022463	Homo sapiens hypothetical protein FLJ14033 similar to hypoxia inducible factor
NM_022462	2 alpha guhunit (HIF-3A) mRNA
ND 6 022461	Homo sapiens hypothetical protein FLJ21939 similar to 5-azacytidine induced
NM_022461	gene 2 (FLJ21939), mRNA
ND4 022452	Homo sapiens ring finger protein 25 (RNF25), mRNA
NM_022453	Homo sapiens likely ortholog of mouse ADP-ribosylation-like factor 6
NM_022374	interacting protein 2 (FLJ23293), mRNA
>D f 000071	Homo sapiens ATP-dependant interferon responsive (ADIR), mRNA
NM_022371	Homo sapiens hypothetical protein FLJ12541 similar to Stra6 (FLJ12541),
NM_022369	mRNA
NM_022367	Homo sapiens hypothetical protein FLJ12287 similar to semaphorins (FLJ12287), mRNA
NM 022359	Homo sapiens similar to rat myomegalin (LOC64182), mRNA
NM 022356	Homo sapiens growth suppressor 1 (GROS1), mRNA
	Homo sapiens spermatogenesis associated 1 (SPATA1), mRNA
NM_022354	Homo sapiens IFRG15 protein (IFRG15), mRNA
	Homo sapiens IFRG15 protein (IFRG15), mRNA  Homo sapiens peptide deformylase-like protein (LOC64146), mRNA

DR 6 000147	1 201 D :
NM_022147	Homo sapiens 28kD interferon responsive protein (IFRG28), mRNA
NM_022140	Homo sapiens erythrocyte protein band 4.1-like 4 (EPB41L4), mRNA
NM_022133	Homo sapiens sorting nexin 16 (SNX16), mRNA
NM_022126	Homo sapiens phospholysine phosphohistidine inorganic pyrophosphate
3 D. C.	phosphatase (LHPP), mRNA
NM_022097	Homo sapiens hepatocellular carcinoma antigen gene 520 (LOC63928), mRNA
NM_022094	Homo sapiens hypothetical protein FLJ20871 similar to FSP27 (FLJ20871), mRNA
NM 022090	Homo sapiens transposon-derived Buster3 transposase-like (LOC63920), mRNA
NM 022074	Homo sapiens hypothetical protein FLJ22794 (FLJ22794), mRNA
NM 022071	Homo sapiens hypothetical protein FLJ20967 (FLJ20967), mRNA
NM 022063	Homo sapiens hypothetical protein FLJ13188 (FLJ13188), mRNA
NM 022060	Homo sapiens hypothetical protein FLJ12816 (FLJ12816), mRNA
NM 022034	Homo sapiens estrogen regulated gene 1 (ERG-1), mRNA
NM 021945	Homo sapiens hypothetical protein FLJ22174 (FLJ22174), mRNA
NM 021944	Homo sapiens hypothetical protein FLJ12154 (FLJ12154), mRNA
NM 021941	Homo sapiens hypothetical protein FLJ21324 (FLJ21324), mRNA
NM 021928	Homo sapiens hypothetical protein FLJ22649 similar to signal peptidase
1.1.1_011>20	SPC22/23 (FLJ22649), mRNA
NM 021927	Homo sapiens hypothetical protein FLJ13220 (FLJ13220), mRNA
NM 021925	Homo sapiens hypothetical protein FLJ21820 (FLJ21820), mRNA
NM 021825	Homo sapiens hypothetical protein MDS025 (MDS025), mRNA
NM 015622	Homo sapiens CGI-43 protein (LOC51622), mRNA
NM 021639	Homo sapiens hypothetical protein SP192 (SP192), mRNA
NM 021637	Homo sapiens hypothetical protein FLJ14084 (FLJ14084), mRNA
NM 021614	Homo sapiens potassium intermediate/small conductance calcium-activated
	channel, subfamily N, member 2 (KCNN2), mRNA
NM 021182	Homo sapiens minor histocompatibility antigen HB-1 (HB-1), mRNA
NM 021170	Homo sapiens bHLH factor Hes4 (LOC57801), mRNA
NM 021146	Homo sapiens angiopoietin-like factor (CDT6), mRNA
NM 005146	Homo sapiens squamous cell carcinoma antigen recognised by T cells (SART1),
	mRNA
NM_021079	Homo sapiens N-myristoyltransferase 1 (NMT1), mRNA
NM_021046	Homo sapiens UHS KerB (LOC57830), mRNA
NM_021018	Homo sapiens H3 histone family, member I (H3FI), mRNA
NM_006643	Homo sapiens serologically defined colon cancer antigen 3 (SDCCAG3), mRNA
NM_017569	Homo sapiens transcription factor (p38 interacting protein) (P38IP), mRNA
NM_015239	Homo sapiens KIAA1035 protein (KIAA1035), mRNA
NM_014977	Homo sapiens KIAA0670 protein/acinus (KIAA0670), mRNA
NM_015176	Homo sapiens KIAA0483 protein (KIAA0483), mRNA
NM_014610	Homo sapiens KIAA0088 protein (KIAA0088), mRNA
NM_015516	Homo sapiens hypothetical protein, estradiol-induced (E2IG4), mRNA
NM_015388	Homo sapiens DKFZP566C243 protein (DKFZP566C243), mRNA
NM_015679	Homo sapiens hypothetical protein (CLONE24922), mRNA
NM_014409	Homo sapiens TAF5-like RNA polymerase II, p300/CBP-associated factor
	(PCAF)-associated factor, 65 kD (TAF5L), mRNA
NM_014368	Homo sapiens LIM homeobox protein 6 (LHX6), mRNA
NM_014315	Homo sapiens host cell factor homolog (LCP), mRNA
NM_012414	Homo sapiens rab3 GTPase-activating protein, non-catalytic subunit (150kD)
	(RAB3-GAP150), mRNA
NM_012219	Homo sapiens muscle RAS oncogene homolog (MRAS), mRNA
NM_007375	Homo sapiens TAR DNA binding protein (TARDBP), mRNA

NB4 007074	Homo sapiens coronin, actin binding protein, 1A (CORO1A), mRNA
NM_007074	Homo sapiens sialyltransferase 4B (beta-galactosidase alpha-2,3-
NM_006927	sialytransferase) (SIAT4B), mRNA
37.6.006061	Homo sapiens RAB35, member RAS oncogene family (RAB35), mRNA
NM_006861	Homo sapiens polymerase (DNA directed), eta (POLH), mRNA
NM_006502	Homo sapiens polymerase (DNA directed), eta (PODI), midva
NM_005710_	Homo sapiens polyglutamine binding protein 1 (PQBP1), mRNA
NM_005168	Homo sapiens ras homolog gene family, member E (ARHE), mRNA
NM_004190	Homo sapiens lipase, gastric (LIPF), mRNA
NM 004132	Homo sapiens hyaluronan binding protein 2 (HABP2), mRNA
NM 004492	Homo sapiens general transcription factor IIA, 2 (12kD subunit) (GTF2A2),
_	mRNA
NM 004824	Homo sapiens chromodomain protein, Y chromosome-like (CDYL), mRNA
NM 003969	Homo sapiens ubiquitin-conjugating enzyme E2M (UBC12 homolog, yeast)
<u> </u>	(JIRE2M) mRNA
NM 002711	Homo saniens protein phosphatase 1, regulatory (inhibitor) subunit 3A (glycogen
	and sarcoplasmic reticulum binding subunit, skeletal muscle) (PPP1R3A),
	mRNA
NM 003847	Homo saniens peroxisomal biogenesis factor 11A (PEX11A), mRNA
NM 002004	Homo sapiens farnesyl diphosphate synthase (farnesyl pyrophosphate synthetase,
14141_002004	dimethylallyltranstransferase, geranyltranstransferase) (FDPS), mRNA
NM 019111	Homo sapiens major histocompatibility complex, class II, DR alpha (HLA-
14141_019111	DRA), mRNA
ND 6 002120	Homo sapiens major histocompatibility complex, class II, DO beta (HLA-DOB),
NM_002120	
37.6 000110	mRNA Homo sapiens major histocompatibility complex, class II, DM beta (HLA-
NM_002118	
	DMB), mRNA
NM_002125	Homo sapiens major histocompatibility complex, class II, DR beta 5 (HLA-DRB5), mRNA
NB4 021092	Homo sapiens major histocompatibility complex, class II, DR beta 4 (HLA-
NM_021983	DRB4), mRNA
ND 4 022555	Homo sapiens major histocompatibility complex, class II, DR beta 3 (HLA-
NM_022555	
27 6 005060	DRB3), mRNA Homo sapiens MAX interacting protein 1 (MXI1), transcript variant 1, mRNA
NM_005962	Homo sapiens MAX interacting protein 1 (MXII), transcript variant 2, mRNA
NM_130439	Homo sapiens MAX interacting protein 1 (MXII), transcript variant 2, mRNA
NM_080923	Homo sapiens protein tyrosine phosphatase, receptor type, C (PTPRC), transcript
	variant 4, mRNA
NM_080922	Homo sapiens protein tyrosine phosphatase, receptor type, C (PTPRC), transcript
	variant 3, mRNA
NM_080921	Homo sapiens protein tyrosine phosphatase, receptor type, C (PTPRC), transcript
_	variant 2, mRNA
NM 130386	Homo sapiens collectin sub-family member 12 (COLEC12), transcript variant I,
-	mRNA
NM 030781	Homo sapiens collectin sub-family member 12 (COLEC12), transcript variant II,
	mRNA
NM 130778	Homo sapiens collagen, type XVII, alpha 1 (COL17A1), transcript variant short,
1414_150776	mRNA
NM 000494	Homo sapiens collagen, type XVII, alpha 1 (COL17A1), transcript variant long,
11111 _000434	mRNA
NIM 001056	Homo sapiens collagen, type XVI, alpha 1 (COL16A1), mRNA
NM_001856	Homo sapiens collagen, type XV, alpha 1 (COL15A1), mRNA
NM_001855	TO THE Sapiens conagen, type Av, alpha I (CODISAL), mich.
NM_058166	Homo sapiens tripartite motif-containing 6 (TRIM6), mRNA
NM_002838	Homo sapiens protein tyrosine phosphatase, receptor type, C (PTPRC), transcript

	variant 1, mRNA
NM 130390	Homo sapiens tripartite motif-containing 34 (TRIM34), transcript variant 3,
112.2_10000	mRNA
NM_130389	Homo sapiens tripartite motif-containing 34 (TRIM34), transcript variant 2,
	mRNA
NM_021616	Homo sapiens tripartite motif-containing 34 (TRIM34), transcript variant 1,
_	mRNA
NM_030950	Homo sapiens ret finger protein (RFP), transcript variant beta, mRNA
NM_130785	Homo sapiens TPTE and PTEN homologous inositol lipid phosphatase (TPIP),
	mRNA
NM_130784	Homo sapiens hypothetical gene supported by AY027807; AY027808
	(LOC93426), mRNA
NM_130783	Homo sapiens similar to neuronal tetraspanin (LOC90139), mRNA
NM_130782	Homo sapiens regulator of G-protein signalling 18 (RGS18), mRNA
NM_130781	Homo sapiens (RAB24), mRNA
NM_130772	Homo sapiens S100Z protein (S100Z), mRNA
NM_130769	Homo sapiens glycoprotein alpha 2 (GPA2), mRNA
NM_130770	Homo sapiens 5-hydroxytryptamine receptor 3 subunit C (HTR3C), mRNA
NM_130768	Homo sapiens GASZ (GASZ), mRNA
NM_130767	Homo sapiens cytosolic acetyl-CoA hydrolase (CACH-1), mRNA
NM_130773	Homo sapiens caspr5 protein (caspr5), mRNA
NM_006510	Homo sapiens ret finger protein (RFP), transcript variant alpha, mRNA
NM_033554	Homo sapiens major histocompatibility complex, class II, DP alpha 1 (HLA-
	DPA1), mRNA
NM_033282	Homo sapiens opsin 4 (melanopsin) (OPN4), mRNA
NM_032035	Homo sapiens MSTP031 protein (MSTP031), mRNA
NM_017882	Homo sapiens ceroid-lipofuscinosis, neuronal 6, late infantile, variant (CLN6),
	mRNA
NM_006983	Homo sapiens matrix metalloproteinase 23B (MMP23B), mRNA
NM_005608	Homo sapiens protein tyrosine phosphatase, receptor type, C-associated protein
	(PTPRCAP), mRNA
NM_004659	Homo sapiens matrix metalloproteinase 23A (MMP23A), mRNA
NM_025091	Homo sapiens hypothetical protein FLJ13330 (FLJ13330), mRNA
NM_130759	Homo sapiens immunity associated protein 1 (IMAP1), mRNA
NM_019841	Homo sapiens transient receptor potential cation channel, subfamily V, member
	5 (TRPV5), mRNA
NM_017584	Homo sapiens aldehyde reductase (aldose reductase) like 6 (ALDRL6), mRNA
NM_017436	Homo sapiens alpha 1,4-galactosyltransferase (A4GALT), mRNA
NM_006480	Homo sapiens regulator of G-protein signalling 14 (RGS14), mRNA
NM_013357	Homo sapiens purine-rich element binding protein G (PURG), mRNA
NM_016155	Homo sapiens matrix metalloproteinase 17 (membrane-inserted) (MMP17),
	mRNA
NM_002813	Homo sapiens proteasome (prosome, macropain) 26S subunit, non-ATPase, 9
	(PSMD9), mRNA
NM_024549	Homo sapiens hypothetical protein FLJ21127 (FLJ21127), mRNA
NM_130441	Homo sapiens dendritic cell lectin b (DLEC), mRNA
NM_015409	Homo sapiens E1A binding protein p400 (EP400), mRNA
NM_003702	Homo sapiens regulator of G-protein signalling 20 (RGS20), mRNA
NM_016113	Homo sapiens transient receptor potential cation channel, subfamily V, member
	2 (TRPV2), mRNA
NM_015530	Homo sapiens likely ortholog of rat golgi stacking protein homolog GRASP55
	(GRASP55), mRNA

NM 005873	Homo sapiens regulator of G-protein signalling 19 (RGS19), mRNA
NM 130469	Homo sapiens Jun dimerization protein 2 (jdp2), mRNA
NM 130468	Homo sapiens dermatan-4-sulfotransferase-1 (D4ST-1), mRNA
NM 130467	Homo sapiens PAGE-5 protein (PAGE-5), mRNA
NM 130463	Homo sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump)
1441_150.05	(ATP6G), mRNA
NM 130459	Homo sapiens torsin family 2, member A (TOR2A), mRNA
NM 021070	Homo sapiens latent transforming growth factor beta binding protein 3 (LTBP3),
14141_021070	mRNA
NM_020865	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 36 (DDX36),
14141_020003	mRNA
NM 016304	Homo sapiens 60S ribosomal protein L30 isolog (LOC51187), mRNA
NM 130443	Homo sapiens dipeptidylpeptidase III (DPP3), transcript variant 2, mRNA
NM 005700	Homo sapiens dipeptidylpeptidase III (DPP3), transcript variant 1, mRNA
NM 018152	Homo sapiens chromosome 20 open reading frame 12 (C20orf12), mRNA
	Homo sapiens exonuclease 1 (EXO1), transcript variant 1, mRNA
NM_006027	Homo sapiens exonuclease 1 (EXO1), transcript variant 1, mRNA  Homo sapiens exonuclease 1 (EXO1), transcript variant 3, mRNA
NM_003686	
NM_130398	Homo sapiens exonuclease 1 (EXO1), transcript variant 2, mRNA  Homo sapiens protein tyrosine phosphatase, receptor type, B (PTPRB), mRNA
NM_002837	Homo sapiens protein tyrosme phosphatase, receptor type, B (F1FRB), mixiA
NM_000775	Homo sapiens cytochrome P450, subfamily IIJ (arachidonic acid epoxygenase)
) The 052056	polypeptide 2 (CYP2J2), mRNA
NM_053056	Homo sapiens cyclin D1 (PRAD1 parathyroid adenomatosis 1) (CCND1),
377 610000	mRNA
NM_012090	Homo sapiens microtubule-actin crosslinking factor 1 (MACF1), transcript
ND 6 017605	variant 1, mRNA
NM_017625	Homo sapiens intelectin (ITLN), mRNA
NM_015839	Homo sapiens ficolin (collagen/fibrinogen domain containing lectin) 2 (hucolin) (FCN2), transcript variant SV3, mRNA
NM_015838	Homo sapiens ficolin (collagen/fibrinogen domain containing lectin) 2 (hucolin)
ND 6 015005	(FCN2), transcript variant SV2, mRNA
NM_015837	Homo sapiens ficolin (collagen/fibrinogen domain containing lectin) 2 (hucolin)
37.5.00000	(FCN2), transcript variant SV1, mRNA
NM_002003	Homo sapiens ficolin (collagen/fibrinogen domain containing) 1 (FCN1), mRNA
NM_016327	Homo sapiens ureidopropionase, beta (UPB1), mRNA
NM_016328	Homo sapiens GTF2I repeat domain containing 1 (GTF2IRD1), transcript
ND4 004100	variant 1, mRNA  Homo sapiens ficolin (collagen/fibrinogen domain containing lectin) 2 (hucolin)
NM_004108	
NIM 002219	(FCN2), transcript variant SV0, mRNA
NM_002318	Homo sapiens lysyl oxidase-like 2 (LOXL2), mRNA  Homo sapiens WNT1 inducible signaling pathway protein 3 (WISP3), transcript
NM_130396	
NTM 002000	variant 2, mRNA  Homo sapiens WNT1 inducible signaling pathway protein 3 (WISP3), transcript
NM_003880	
NTM 002001	variant 1, mRNA
NM_003881	Homo sapiens WNT1 inducible signaling pathway protein 2 (WISP2), mRNA
NM_080838	Homo sapiens WNT1 inducible signaling pathway protein 1 (WISP1), transcript variant 2, mRNA
NM 003882	Homo sapiens WNT1 inducible signaling pathway protein 1 (WISP1), transcript
	variant 1, mRNA
NM_000651	Homo sapiens complement component (3b/4b) receptor 1, including Knops
	blood group system (CR1), transcript variant S, mRNA
NM_000573	Homo sapiens complement component (3b/4b) receptor 1, including Knops
	blood group system (CR1), transcript variant F, mRNA
_	

NM_006069 NM_130385	Homo sapiens murine retrovirus integration site 1 homolog (MRVII), transcript variant 1, mRNA
NM_130385	
	Homo sapiens murine retrovirus integration site 1 homolog (MRVI1), transcript variant 2, mRNA
NM 018492	Homo sapiens T-LAK cell-originated protein kinase (TOPK), mRNA
NM 002462	Homo sapiens myxovirus (influenza virus) resistance 1, interferon-inducible
_	protein p78 (mouse) (MX1), mRNA
NM_015920	Homo sapiens ribosomal protein S27-like (RPS27L), mRNA
NM_016183	Homo sapiens ribosomal protein, large, P0-like (RPLP0L), mRNA
NM_080746	Homo sapiens ribosomal protein L10-like (RPL10L), mRNA
NM_032236	Homo sapiens FLJ23277 protein (FLJ23277), mRNA
NM_032784	Homo sapiens thrombospondin (FLJ14440), mRNA
NM_080731	Homo sapiens intermediate filament-like MGC:2625 (DKFZP586I2223).
	transcript variant 3, mRNA
NM_080730	Homo sapiens intermediate filament-like MGC:2625 (DKFZP586I2223),
	transcript variant 2, mRNA
NM_015945	Homo sapiens ovarian cancer overexpressed 1 (OVCOV1), mRNA
NM_018018	Homo sapiens solute carrier family 38, member 4 (SLC38A4), mRNA
NM_022451	Homo sapiens AD24 protein (AD24), mRNA
NM_020830	Homo sapiens phosphoinositide-binding protein SR1 (FENS-1), mRNA
NM_033630	Homo sapiens SCAN domain containing 1 (SCAND1), transcript variant 2, mRNA
NM_016558	Homo sapiens SCAN domain containing 1 (SCAND1), transcript variant 1, mRNA
NM_015438	Homo sapiens intermediate filament-like MGC:2625 (DKFZP586I2223),
	transcript variant 1, mRNA
NM_007371	Homo sapiens bromodomain containing 3 (BRD3), mRNA
NM_005104	Homo sapiens bromodomain containing 2 (BRD2), mRNA
NM_005031	Homo sapiens FXYD domain containing ion transport regulator 1
	(phospholemman) (FXYD1), transcript variant a, mRNA
NM_021902	Homo sapiens FXYD domain containing ion transport regulator 1
	(phospholemman) (FXYD1), transcript variant b, mRNA
NM_014164	Homo sapiens FXYD domain-containing ion transport regulator 5 (FXYD5), mRNA
NM_002463	Homo sapiens myxovirus (influenza virus) resistance 2 (mouse) (MX2), mRNA
NM_014577	Homo sapiens bromodomain containing 1 (BRD1), mRNA
NM_021004	Homo sapiens peroxisomal short-chain alcohol dehydrogenase (humNRDR), mRNA
NM_020399	Homo sapiens PDZ/coiled-coil domain binding partner for the rho-family GTPase TC10 (PIST), mRNA
NM_017935	Homo sapiens hypothetical protein FLJ20706 (BANK), mRNA
NM_018244	Homo sapiens chromosome 20 open reading frame 44 (C20orf44), mRNA
NM_016100	Homo sapiens N-acetyltransferase 5 (ARD1 homolog, S. cerevisiae) (NAT5), mRNA
NM_016045	Homo sapiens chromosome 20 open reading frame 45 (C20orf45), mRNA
NM_007363	Homo sapiens non-POU domain containing, octamer-binding (NONO), mRNA
NM_002438	Homo sapiens mannose receptor, C type 1 (MRC1), mRNA
NM_015092	Homo sapiens PI-3-kinase-related kinase SMG-1 (SMG1), mRNA
NM_018993	Homo sapiens RAB5 interacting protein 2 (RIN2), mRNA
NM_080841	Homo sapiens protein tyrosine phosphatase, receptor type, A (PTPRA),
	transcript variant 3, mRNA
NM 080840	Homo sapiens protein tyrosine phosphatase, receptor type, A (PTPRA),

	transcript variant 2, mRNA
NM_002836	Homo sapiens protein tyrosine phosphatase, receptor type, A (PTPRA),
	transcript variant 1, mRNA
NM_024832	Homo sapiens RAB5 interacting protein 3 (RIN3), mRNA
NM_023915	Homo sapiens G protein-coupled receptor 87 (GPR87), mRNA
NM_003029	Homo sapiens SHC (Src homology 2 domain containing) transforming protein 1
	(SHC1), mRNA
NM_018490	Homo sapiens G protein-coupled receptor 48 (GPR48), mRNA
NM_016020	Homo sapiens homolog of yeast mitochondrial transcription factor B (mtTFB),
	MRNA
NM_014475	Homo sapiens dihydrodiol dehydrogenase (dimeric) (DHDH), mRNA
NM_006065	Homo sapiens signal-regulatory protein beta 1 (SIRPB1), mRNA
NM_005527	Homo sapiens heat shock 70kD protein 1-like (HSPA1L), mRNA
NM_004648	Homo sapiens protein tyrosine phosphatase, non-receptor type substrate 1
	(PTPNS1), mRNA
NM_004480	Homo sapiens fucosyltransferase 8 (alpha (1,6) fucosyltransferase) (FUT8),
	mRNA
NM_003667	Homo sapiens G protein-coupled receptor 49 (GPR49), mRNA
NM_130434	Homo sapiens dipeptidylpeptidase 8 (DPP8), transcript variant 1, mRNA
NM_017743	Homo sapiens dipeptidylpeptidase 8 (DPP8), transcript variant 2, mRNA
NM_002122	Homo sapiens major histocompatibility complex, class II, DQ alpha 1 (HLA-
	DQA1), mRNA
NM_006442	Homo sapiens DR1-associated protein 1 (negative cofactor 2 alpha) (DRAP1),
	mRNA
NM_080918	Homo sapiens deoxyguanosine kinase (DGUOK), transcript variant 2, nuclear
	gene encoding mitochondrial protein, mRNA
NM_080917	Homo sapiens deoxyguanosine kinase (DGUOK), transcript variant 3, nuclear
	gene encoding mitochondrial protein, mRNA
NM_080916	Homo sapiens deoxyguanosine kinase (DGUOK), transcript variant 1, nuclear
	gene encoding mitochondrial protein, mRNA
NM_080915	Homo sapiens deoxyguanosine kinase (DGUOK), transcript variant 5, nuclear
	gene encoding mitochondrial protein, mRNA
NM_001929	Homo sapiens deoxyguanosine kinase (DGUOK), transcript variant 4 nuclear
	gene encoding mitochondrial protein, mRNA
NM_080815	Homo sapiens collagen, type XIII, alpha 1 (COL13A1), transcript variant 19,
	mRNA
NM_080814	Homo sapiens collagen, type XIII, alpha 1 (COL13A1), transcript variant 18,
	mrna
NM_080813	Homo sapiens collagen, type XIII, alpha 1 (COL13A1), transcript variant 17,
	mRNA
NM_080812	Homo sapiens collagen, type XIII, alpha 1 (COL13A1), transcript variant 16,
	mRNA
NM_080811	Homo sapiens collagen, type XIII, alpha 1 (COL13A1), transcript variant 15,
	mRNA
NM_080810	Homo sapiens collagen, type XIII, alpha 1 (COL13A1), transcript variant 14,
	mRNA
NM_080809	Homo sapiens collagen, type XIII, alpha 1 (COL13A1), transcript variant 13,
	mRNA
NM_080808	Homo sapiens collagen, type XIII, alpha 1 (COL13A1), transcript variant 12,
	mRNA
NM_080807	Homo sapiens collagen, type XIII, alpha 1 (COL13A1), transcript variant 11,
	mRNA

NM_080806	Homo sapiens collagen, type XIII, alpha 1 (COL13A1), transcript variant 10, mRNA
NM_080805	Homo sapiens collagen, type XIII, alpha 1 (COL13A1), transcript variant 9, mRNA
NM_080804	Homo sapiens collagen, type XIII, alpha 1 (COL13A1), transcript variant 8, mRNA
NM_080803	Homo sapiens collagen, type XIII, alpha 1 (COL13A1), transcript variant 7, mRNA
NM_080802	Homo sapiens collagen, type XIII, alpha 1 (COL13A1), transcript variant 6, mRNA
NM_080801	Homo sapiens collagen, type XIII, alpha 1 (COL13A1), transcript variant 5, mRNA
NM_080800	Homo sapiens collagen, type XIII, alpha 1 (COL13A1), transcript variant 4, mRNA
NM_080799	Homo sapiens collagen, type XIII, alpha 1 (COL13A1), transcript variant 3, mRNA
NM_080798	Homo sapiens collagen, type XIII, alpha 1 (COL13A1), transcript variant 2, mRNA
NM_005203	Homo sapiens collagen, type XIII, alpha 1 (COL13A1), transcript variant 1, mRNA
NM_004395	Homo sapiens drebrin 1 (DBN1), transcript variant 1, mRNA
NM_080881	Homo sapiens drebrin 1 (DBN1), transcript variant 2, mRNA
NM_080792	Homo sapiens brain-immunoglobulin-like molecule with tyrosine-based activation motifs (BIT), mRNA
NM_080816	Homo sapiens signal-regulatory protein beta 2 (SIRPB2), transcript variant 2, mRNA
NM_018556	Homo sapiens signal-regulatory protein beta 2 (SIRPB2), transcript variant 1, mRNA
NM_000787	Homo sapiens dopamine beta-hydroxylase (dopamine beta-monooxygenase) (DBH), mRNA
NM_080426	Homo sapiens GNAS complex locus (GNAS), transcript variant 2, mRNA
NM_080425	Homo sapiens GNAS complex locus (GNAS), transcript variant 3, mRNA
NM_000516	Homo sapiens GNAS complex locus (GNAS), transcript variant 1, mRNA
NM_006571	Homo sapiens novel RGD-containing protein (WS-3), mRNA
NM_080926	Homo sapiens hypothetical protein similar to KIAA0187 gene product (LOC96610), mRNA
NM_080924	Homo sapiens hypothetical protein similar to CGI-67 protein (LOC91219), mRNA
NM_080925	Homo sapiens hypothetical protein similar to topoisomerase (DNA) III beta (H. sapiens) (LOC129020), mRNA
NM_080914	Homo sapiens asialoglycoprotein receptor 2 (ASGR2), transcript variant 3, mRNA
NM_080913	Homo sapiens asialoglycoprotein receptor 2 (ASGR2), transcript variant 2, mRNA
NM_080912	Homo sapiens asialoglycoprotein receptor 2 (ASGR2), transcript variant H2', mRNA
NM_001181	Homo sapiens asialoglycoprotein receptor 2 (ASGR2), transcript variant 1, mRNA
NM_001671	Homo sapiens asialoglycoprotein receptor 1 (ASGR1), mRNA
NM_005065	Homo sapiens sel-1 suppressor of lin-12-like (C. elegans) (SEL1L), mRNA
NM_014978	Homo sapiens VPS10 domain receptor protein SORCS 3 (SORCS3), mRNA
NM_015230	Homo sapiens centaurin, delta 1 (CENTD1), mRNA

Ĭ.Š

NT 050050	
NM_052868	Homo sapiens immunoglobulin superfamily, member 8 (IGSF8), mRNA
NM_032782	Homo sapiens hypothetical protein FLJ14428 (TIM3) mRNA
NM_032309	Homo sapiens chromosome 2 open reading frame 9 (C2orf9) mPNA
NM_021625	Homo sapiens transient receptor potential cation channel subfamily V momber
377	14 (1RF v4), mRNA
NM 020960	Homo sapiens G protein-coupled receptor 107 (GPR107), mRNA
NM_024503	Homo sapiens human immunodeficiency virus type I enhancer hinding protein 3
NM 024112	(III V EF 3), INKINA
NM_015192	Homo sapiens chromosome 9 open reading frame 16 (C9orf16), mRNA
	mRNA phospholipase C, beta 1 (phosphoinositide-specific) (PLCB1),
NM_022481	Homo sapiens ARF-GAP, RHO-GAP, ankyrin repeat and plekstrin homology
	domains-containing protein 3 (ARAP3), mRNA
NM_021634	Homo sapiens leucine-rich repeat-containing G protein-coupled receptor 7
	(LGR7), mRNA
NM_013305	Homo sapiens sialyltransferase 8E (alpha-2, 8-polysialytransferase) (SIAT8E),
	mRNA (SIAT8E),
NM_019069	Homo sapiens WD repeat domain 5B (WDR5B), mRNA
NM 016179	Homo sapiens transient receptor potential cation channel, subfamily C, member
	4 (TRPC4), mRNA
NM 016592	Homo saniens GNAS complex logge (CNAS)
NM 014007	Homo sapiens GNAS complex locus (GNAS), transcript variant 4, mRNA Homo sapiens zinc finger protein 297B (ZNF297B), mRNA
NM_012471	Homo sapiens transient recent with the sapient recent
	Homo sapiens transient receptor potential cation channel, subfamily C, member 5 (TRPC5), mRNA
NM_012459	Homo caniena translaces of:
- 1212 133	Homo sapiens translocase of inner mitochondrial membrane 8 homolog B (yeast) (TIMM8B), mRNA
NM 004621	Home conjugate
144_004021	Homo sapiens transient receptor potential cation channel, subfamily C, member
NM 003304	10 (11d CO), IIICIVA
- 1212_00550-4	Homo sapiens transient receptor potential cation channel, subfamily C, member 1 (TRPC1), mRNA
NM 002124	Home conjugation 1:
1.1.1_002124	Homo sapiens major histocompatibility complex, class II, DR beta 1 (HLA-
NM 000972	LDKD1), IIIKNA
NM_130384	Homo sapiens ribosomal protein L7a (RPL7A), mRNA
.1012_130364	Homo sapiens three prime repair exonuclease 1 (TREX1), transcript variant 6, mRNA
VM_033627	Homo sapiens three prime repair exonuclease 1 (TREX1), transcript variant 2,
	mRNA mRNA
VM_032166	Homo sapiens three prime repair exonuclease 1 (TREX1), transcript variant 5,
·-	mRNA
IM_024996	Homo sapiens mitochondrial elongation factor G (EFG1), mRNA
IM 033629	Homo sapiens three prime repair examples and (EFGI), mRNA
	Homo sapiens three prime repair exonuclease 1 (TREX1), transcript variant 4, mRNA
VM_033628	
	Homo sapiens three prime repair exonuclease 1 (TREX1), transcript variant 3, mRNA
M 016381	
	Homo sapiens three prime repair exonuclease 1 (TREX1), transcript variant 1, mRNA
IM_031892	ILLUA
	Homo sapiens SH3-domain kinase binding protein 1 (SH3KBP1), mRNA
111_005500	monio sapiens N-acetyltransferase 8 (camello like) (NATS) mpNA
m_071033	Homo sapiens peptide YY, 2 (seminalplasmin) (PYY2) mRNA
M 021002	
WI_021092	nomo sapiens pancreatic polypeptide 2 (PPY2) mRNA
M_021190	Homo sapiens pancreatic polypeptide 2 (PPY2), mRNA  Homo sapiens polypyrimidine tract binding protein 2 (PTBP2), mRNA  Homo sapiens tachykinin, precursor 1 (substance K, substance P, neurokinin 1,

	neurokinin 2, neuromedin L, neurokinin alpha, neuropeptide K, neuropeptide gamma) (TAC1), transcript variant delta, mRNA
NM_013997	Homo sapiens tachykinin, precursor 1 (substance K, substance P, neurokinin 1,
1414_013997	neurokinin 2, neuromedin L, neurokinin alpha, neuropeptide K, neuropeptide
	gamma) (TAC1), transcript variant gamma, mRNA
NM 013996	
14141_013990	Homo sapiens tachykinin, precursor 1 (substance K, substance P, neurokinin 1,
	neurokinin 2, neuromedin L, neurokinin alpha, neuropeptide K, neuropeptide
ND ( 01/025	gamma) (TAC1), transcript variant alpha, mRNA
NM_016235	Homo sapiens G protein-coupled receptor, family C, group 1, member B (GPRC5B), mRNA
NM 004630	
NM 000230	Homo sapiens splicing factor 1 (SF1), mRNA
	Homo sapiens leptin (obesity homolog, mouse) (LEP), mRNA
NM_003185	Homo sapiens TAF4 RNA polymerase II, TATA box binding protein (TBP)-
NR 6 002100	associated factor, 135 kD (TAF4), mRNA
NM_003182	Homo sapiens tachykinin, precursor 1 (substance K, substance P, neurokinin 1,
	neurokinin 2, neuromedin L, neurokinin alpha, neuropeptide K, neuropeptide
) T 6 000550	gamma) (TAC1), transcript variant beta, mRNA
NM_002772	Homo sapiens protease, serine, 7 (enterokinase) (PRSS7), mRNA
NM_005857	Homo sapiens zinc metalloproteinase (STE24 homolog, yeast) (ZMPSTE24),
	mRNA
NM_006103	Homo sapiens WAP four-disulfide core domain 2 (WFDC2), transcript variant 1,
	mRNA
NM_080736	Homo sapiens WAP four-disulfide core domain 2 (WFDC2), transcript variant 2,
	mRNA
NM_080735	Homo sapiens WAP four-disulfide core domain 2 (WFDC2), transcript variant 5, mRNA
NM_080734	Homo sapiens WAP four-disulfide core domain 2 (WFDC2), transcript variant 4, mRNA
NM_080733	Homo sapiens WAP four-disulfide core domain 2 (WFDC2), transcript variant 3,
	mRNA
NM_021197	Homo sapiens WAP four-disulfide core domain 1 (WFDC1), mRNA
NM_007128	Homo sapiens pre-B lymphocyte gene 1 (VPREB1), mRNA
NM_006373	Homo sapiens vesicle amine transport protein 1 (VATI), mRNA
NM_003105	Homo sapiens sortilin-related receptor, L(DLR class) A repeats-containing (SORL1), mRNA
NM_020777	Homo sapiens VPS10 domain receptor protein (SORCS2), mRNA
NM_052918	Homo sepiens VPS10 domain receptor protein (SORCS2), mRNA
NM 022553	Homo sapiens VPS10 domain receptor protein SORCS 1 (SORCS1), mRNA
11111_022333	Homo sapiens SAC2 suppressor of actin mutations 2-like (yeast) (SACM2L),
NM 004843	transcript variant 2, mRNA
	Homo sapiens class I cytokine receptor (WSX1), mRNA
NM_080564	Homo sapiens SAC2 suppressor of actin mutations 2-like (yeast) (SACM2L), transcript variant 1, mRNA
NM 006711	Homo sapiens RNA binding protein S1, serine-rich domain (RNPS1), transcript
14141_000711	variant 1, mRNA
NM 080594	
14141_000334	Homo sapiens RNA binding protein S1, serine-rich domain (RNPS1), transcript variant 2, mRNA
NM 100486	Homo sapiens WW domain-containing adapter with a coiled-coil region (WAC),
	transcript variant 3, mRNA
NM 100264	Homo sapiens WW domain-containing adapter with a coiled-coil region (WAC),
·	transcript variant 2, mRNA
NM_016628	Homo sapiens WW domain-containing adapter with a coiled-coil region (WAC),
10020	transcript variant 1, mRNA
L <u></u>	Larrange A, MILLANI

77.5 00.5	
NM_005701	The state of the s
NM_014810	Homo sapiens centrosome-associated protein 350 (CAP350) mPNA
NM_013325	Homo sapiens KIAA0943 protein (Apg4B), mRNA
NM_020235	Homo sapiens bobby sox homolog (Drosophila) (BRY) mpNA
NM_019118	Homo sapiens hypothetical protein RP4-6221.5 (RP4-6221.5) mPNA
NM_016312	Tionio sapiens w w domain binding protein 11 (WRP11) mpNIA
NM_018706	Homo sapiens KIAA1630 protein (KIAA1630) mRNA
NM_080599	Homo sapiens regulator of nonsense transcripts 2 (RENT2), transcript variant 1,
NM_015542	Homo sapiens regulator of nonsense transcripts 2 (RENT2), transcript variant 2,
NM_002911	Homo sapiens regulator of nonsense transcripts 1 (RENT1), mRNA
NM_002833	fromo sapiens protein tyrosine phosphatase, non-recentor type 9 (PTPNO)
NM_080589	Homo sapiens protein tyrosine phosphatase, non-receptor type 7 (PTPN7),
	tanscript variant 5, mknA
NM_080588	Homo sapiens protein tyrosine phosphatase, non-receptor type 7 (PTPN7),
	transcript variant 2. mknA
NM_002832	Homo sapiens protein tyrosine phosphatase, non-receptor type 7 (PTPN7),
) T. C.	
NM_007039	Homo sapiens protein tyrosine phosphatase, non-receptor type 21 (PTPN21),
ND4 014060	India
NM_014369	Homo sapiens protein tyrosine phosphatase, non-receptor type 18 (brain-derived)
NTM 005401	
NM_005401	Homo sapiens protein tyrosine phosphatase, non-receptor type 14 (PTPN14),
NM_002835	HIII 427
14141_002833	Homo sapiens protein tyrosine phosphatase, non-receptor type 12 (PTPN12),
NM_080685	1 mm 4V
14147_080083	Homo sapiens protein tyrosine phosphatase, non-receptor type 13 (APO-1/CD95
NM_080684	1 (1 ds) -dssociated phosphatase) (P   PN   3) franscript variant 4 mDNA
11111_000004	Tionio sapiens protein tyrosine phosphatase, non recentor time 12 (A DO 1 (OD 0)
NM_080683	1 (1 ds) dssociated phosphalase) (P1PN13) transcript vorient 2 DNIA
	Homo sapiens protein tyrosine phosphatase, non-receptor type 13 (APO-1/CD95
NM_080601	1 (2 db) dbsootated phosphalase (PTPN13) transcript voriont 1 mpN14
	Homo sapiens protein tyrosine phosphatase, non-receptor type 11 (PTPN11), transcript variant 2, mRNA
NM 002834	Homo saniens protein turosino pla milita
_	Homo sapiens protein tyrosine phosphatase, non-receptor type 11 (PTPN11), transcript variant 1, mRNA
NM_006399	Homo saniens hasic leucine zinner transcription Co. A.
_	Homo sapiens basic leucine zipper transcription factor, ATF-like (BATF), mRNA
NM_006709	Homo sapiens HLA-B associated transcript 8 (BAT8), transcript variant
	NG36/G9a, mRNA
NM_033177	Homo sapiens HLA-B associated transcript 4 (BAT4), mRNA
NM_004639	Homo sapiens HLA-B associated transcript 3 (BAT3), transcript variant 1,
	MICHA
NM_080703	Homo sapiens HLA-B associated transcript 3 (BAT3), transcript variant 3,
	mulli
NM_080702	Homo sapiens HLA-B associated transcript 3 (BAT3), transcript variant 2,
	THE WILL
NM_004638	Homo sapiens HLA-B associated transcript 2 (BAT2), transcript variant 2,
	AMC (1)
NM_080686	Homo sapiens HLA-B associated transcript 2 (BAT2), transcript variant 1,
_	tanscript 2 (DA12), transcript variant 1,

	DNA
NM 004640	mRNA Homo sapiens HLA-B associated transcript 1 (BAT1), transcript variant 1,
11111_004640	mRNA
NM 080598	Homo sapiens HLA-B associated transcript 1 (BAT1), transcript variant 2,
1117_00000	mRNA
NM 080797	Homo sapiens death associated transcription factor 1 (DATF1), transcript variant
1	3, mRNA
NM_080796	Homo sapiens death associated transcription factor 1 (DATF1), transcript variant
	2, mRNA
NM_022105	Homo sapiens death associated transcription factor 1 (DATF1), transcript variant
37 6 001000	1, mRNA
NM_021080	Homo sapiens disabled homolog 1 (Drosophila) (DAB1), mRNA Homo sapiens dachshund homolog (Drosophila) (DACH), transcript variant 2,
NM_080760	mRNA
NM 080759	Homo sapiens dachshund homolog (Drosophila) (DACH), transcript variant 1,
1111_000757	mRNA
NM_004392	Homo sapiens dachshund homolog (Drosophila) (DACH), transcript variant 3,
	mRNA
NM_005996	Homo sapiens T-box 3 (ulnar mammary syndrome) (TBX3), transcript variant 1,
	mRNA
NM_016569	Homo sapiens T-box 3 (ulnar mammary syndrome) (TBX3), transcript variant 2,
	mRNA
NM_016954	Homo sapiens T-box 22 (TBX22), mRNA
NM_080701	Homo sapiens three prime repair exonuclease 2 (TREX2), transcript variant 4, mRNA
NM_080700	Homo sapiens three prime repair exonuclease 2 (TREX2), transcript variant 3,
	mRNA
NM_080699	Homo sapiens three prime repair exonuclease 2 (TREX2), transcript variant 2,
37.5 017510	mRNA
NM_017518	Homo sapiens three prime repair exonuclease 2 (TREX2), transcript variant 5,
NM_007205	mRNA Homo sapiens three prime repair exonuclease 2 (TREX2), transcript variant 1,
NM_007203	mRNA
NM_080632	Homo sapiens similar to yeast Upf3, variant B (UPF3B), transcript variant 1,
1.1.12_00000	mRNA
NM_023010	Homo sapiens similar to yeast Upf3, variant B (UPF3B), transcript variant 2,
	mRNA
NM_080687	Homo sapiens similar to yeast Upf3, variant A (UPF3A), transcript variant 2,
	mRNA
NM_023011	Homo sapiens similar to yeast Upf3, variant A (UPF3A), transcript variant 1,
77.5 000.600	mRNA
NM_080630	Homo sapiens collagen, type XI, alpha 1 (COL11A1), transcript variant C, mRNA
NM_080629	Homo sapiens collagen, type XI, alpha 1 (COL11A1), transcript variant B,
NW_080029	mRNA
NM 001854	Homo sapiens collagen, type XI, alpha 1 (COL11A1), transcript variant A,
	mRNA
NM_080791	Homo sapiens acid phosphatase, testicular (ACPT), transcript variant A3, mRNA
NM_001639	Homo sapiens amyloid P component, serum (APCS), mRNA
NM_080790	Homo sapiens acid phosphatase, testicular (ACPT), transcript variant A2, mRNA
NM_080789	Homo sapiens acid phosphatase, testicular (ACPT), transcript variant A1, mRNA
NM 033068	Homo sapiens acid phosphatase, testicular (ACPT), transcript variant A, mRNA

177	PCT/US03/0502
NM_00164	
NM_01448	
	gene encoding mitochondrial much endonuclease-like 2 (APEXL2), nuclear
NM_08064	Homo sapiens APEX nuclease (multifunctional DNA repair enzyme) (APEX),
	transcript variant 3, mRNA
NM_08064	8 Homo saniens APEY pyels a control of the saniens APEY pyels a control of the saniens aperts and the saniens are saniens and the saniens are saniens and the saniens are sani
	8 Homo sapiens APEX nuclease (multifunctional DNA repair enzyme) (APEX), transcript variant 2, mRNA
NM_00164	1 Homo saniens A PEV much
	Homo sapiens APEX nuclease (multifunctional DNA repair enzyme) (APEX), transcript variant 1, mRNA
NM 080839	Homo saniens similar to
NM_080927	Homo sapiens similar to gamma-glutamyltransferase 1 (LOC91227), mRNA Homo sapiens endothelial and smooth muscle cell deciral.
	Drotein (FSDN) DNIA Indisore cell-derived neuropilin-like
NM_030969	Homo saniens hymothetical
NM 080920	
NM_021168	Homo sapiens gamma-glutamyltransferase-like activity 4 (GGTLA4), mRNA Homo sapiens RAR (RAS like GTPASE) like (RABL)
NM_080842	Homo sapiens RAR (RAS like GTPASE) like (RARL), mRNA Homo sapiens hypothetical gene similar
	Homo sapiens hypothetical gene similar to gamma-glutamyltransferase-like activity 1 (LOC129026), mRNA
NM_031460	activity 1 (LOC129026), mRNA
	Homo sapiens potassium channel and control and control
NM 033056	(KCNK17), mRNA
NM_053283	The profit of the state of the
NM 033518	
NM_021160	Tionio Sapiens solute carrier family 20
NM_002279	Homo sapiens HLA-B associated transcript 5 (BAT5), mRNA  Homo sapiens keratin hair acidio 3D (KDR)
NM_004138	Homo sapiens keratin, hair, acidic, 3B (KRTHA3B), mRNA  Homo sapiens keratin hair acidic, 3R (KRTHA3B), mRNA
NM_016310	Homo sapiens keratin, hair, acidic, 3B (KRTHA3B), mRNA Homo sapiens polymerase (RNA) III (RNA)
14141_010310	Homo sapiens polymerase (RNA) III (DNA directed) polypeptide K (12.3 kD)
NM_031991	(POLR3K), mRNA (DVA directed) polypeptide K (12.3 kD)
14141_021991	Homo sapiens polypyrimidine tract hinding protein 1 (Dans vi
VM_031990	Homo sapiens polypyrimidine tract binding protein 1 (PTBP1), transcript variant
MM_031990	Homo sapiens polypyrimidine tract binding protein 1 (PTBP1), transcript variant 1, mRNA
VM_002819	2, mRNA (PTBP1), transcript variant
VIVI_002819	Homo sapiens polypyrimidine tract hinding protein 1 (Proposition)
IM_030930	Homo sapiens polypyrimidine tract binding protein 1 (PTBP1), transcript variant 1, mRNA
DM 022454	Homo saniens unc. 02 homes. Discontinuity
M_022454	Homo sapiens SRY-related HMG-box transcription factor SOX17 (SOX17),
14 004655	mRNA and an action sold (SOX17)
M_004652	Homo sapiens ubiquitin engais
N	Drosophila) (USP9X), transcript variant 1, mRNA
M_021906	Homo sapiens ubiquitin enceif
	Homo sapiens ubiquitin specific protease 9, X chromosome (fat facets-like Homo sapiens membrane variant 2, mRNA
M_022349	Homo sapiens membrane-spanning 4.4
	Homo sapiens membrane-spanning 4-domains, subfamily A, member 6A (MS4A6A), mRNA
V1_U2Z122	Homo sapiens matrix metallarmet
M_006387	Homo sapiens calcium homeostasis endoplasmic reticulum protein (CHERP),
	mRNA monicostasis endoplasmic reticulum protein (CHERP)
1_006918	Homo sapiens sterol C5 docet
	Homo sapiens sterol-C5-desaturase (ERG3 delta-5-desaturase homolog, fungal)-
1_020151	Homo sapiens STADT 1
	Homo sapiens START domain containing 7 (STARD7), mRNA
I_013351   I	Homo sapiens solute carrier family 38, member 2 (SLC38A2), mRNA Homo sapiens T-box 21 (TBX21), mRNA
	Danie Suprems I-DOX 71 (TRY21) DATA
006993 T	Jomo sanione and LI (IDAZI), MKNA
L_006993 F	Homo sapiens T-box 21 (TBX21), mRNA Homo sapiens nucleophosmin/nucleoplasmin, 3 (NPM3), mRNA Homo sapiens transient receptor potential cation channel, subfamily M, member

	1 (TDDM1) mDNA
ND4 007244	1 (TRPM1), mRNA
NM_007244	Homo sapiens proline rich 4 (lacrimal) (PROL4), mRNA
NM_006758	Homo sapiens U2(RNU2) small nuclear RNA auxillary factor 1 (U2AF1), mRNA
NM_006264	Homo sapiens protein tyrosine phosphatase, non-receptor type 13 (APO-1/CD95
	(Fas)-associated phosphatase) (PTPN13), transcript variant 2, mRNA
NM_006055	Homo sapiens LanC lantibiotic synthetase component C-like 1 (bacterial) (LANCL1), mRNA
NM_005716	Homo sapiens regulator of G-protein signalling 19 interacting protein 1 (RGS19IP1), mRNA
NM_005149	Homo sapiens T-box 19 (TBX19), mRNA
NM 004231	Homo sapiens ATPase, vacuolar, 14 kD (ATP6S14), mRNA
NM_000275	Homo sapiens oculocutaneous albinism II (pink-eye dilution homolog, mouse)
	(OCA2), mRNA
NM_001384	Homo sapiens diptheria toxin resistance protein required for diphthamide biosynthesis-like 2 (S. cerevisiae) (DPH2L2), mRNA
NM_000062	Homo sapiens serine (or cysteine) proteinase inhibitor, clade G (C1 inhibitor),
1111_000002	member 1, (angioedema, hereditary) (SERPING1), mRNA
NM_003307	Homo sapiens transient receptor potential cation channel, subfamily M, member 2 (TRPM2), mRNA
NM_003807	Homo sapiens tumor necrosis factor (ligand) superfamily, member 14 (TNFSF14), mRNA
NM 002984	Homo sapiens small inducible cytokine A4 (SCYA4), mRNA
NM 002105	Homo sapiens H2A histone family, member X (H2AFX), mRNA
NM 005331	Homo sapiens hemoglobin, theta 1 (HBQ1), mRNA
NM 000558	Homo sapiens hemoglobin, alpha 1 (HBA1), mRNA
NM 000517	Homo sapiens hemoglobin, alpha 2 (HBA2), mRNA
NM 012262	Homo sapiens heparan sulfate 2-O-sulfotransferase 1 (HS2ST1), mRNA
NM 021213	Homo sapiens phosphatidylcholine transfer protein (PCTP), mRNA
NM 018960	Homo sapiens glycine N-methyltransferase (GNMT), mRNA
NM 017807	Homo sapiens O-sialoglycoprotein endopeptidase (OSGEP), mRNA
NM_016732	Homo sapiens RNA binding protein (autoantigenic, hnRNP-associated with
	lethal yellow) (RALY), transcript variant 1, mRNA
NM_014483	Homo sapiens RNA binding motif, single stranded interacting protein (RBMS3), mRNA
NM_012320	Homo sapiens lysophospholipase 3 (LYPLA3), mRNA
NM_000184	Homo sapiens hemoglobin, gamma G (HBG2), mRNA
NM_005330	Homo sapiens hemoglobin, epsilon 1 (HBE1), mRNA
NM_007367	Homo sapiens RNA binding protein (autoantigenic, hnRNP-associated with lethal yellow) (RALY), transcript variant 2, mRNA
NM 005332	Homo sapiens hemoglobin, zeta (HBZ), mRNA
NM 005438	Homo sapiens FOS-like antigen 1 (FOSL1), mRNA
NM 000158	Homo sapiens glucan (1,4-alpha-), branching enzyme 1 (glycogen branching
	enzyme, Andersen disease, glycogen storage disease type IV) (GBE1), mRNA
NM_000559	Homo sapiens hemoglobin, gamma A (HBG1), mRNA
NG_000007	Homo sapiens genomic beta globin region (HBB@) on chromosome 11
NG_000006	Homo sapiens genomic alpha globin region (HBA@) on chromosome 16
NM_030964	Homo sapiens sprouty homolog 4 (Drosophila) (SPRY4), mRNA
NM_021181	Homo sapiens 19A24 protein (CRACC), mRNA
NM_004654	Homo sapiens ubiquitin specific protease 9, Y chromosome (fat facets-like Drosophila) (USP9Y), mRNA
NM_018518	Homo sapiens MCM10 minichromosome maintenance deficient 10 (S.

	PCT/US03/05
NM_0185	cerevisiae) (MCM10), mRNA
- 11/1_0103	FIGURE Sapiens solute commission C. 11 4 5 5
NM_0182	member 10 (SLC16A10), mRNA
NM_0160	Tionio sapiens kin of IDDE 111- (D.
NM_0068	Homo sapiens chromosome 20 open reading frame 9 (C20orf9), mRNA  Homo sapiens solute carrier family 38, marghes 2 (C20orf9), mRNA
NM_0037	Homo sapiens solute carrier family 38, member 3 (SLC38A3), mRNA Homo sapiens oxidative 3 alpha hydroxystarsid 1
1/1/1/_0037	Homo sapiens oxidative 3 alamy 50, member 3 (SLC38A3), mRNA
NC 0000	dehydrogenase: 3-bydroyysteroid dehydrogenase: retinol
NG_00000	Homo sapiens genomic small history (RODH), mRNA
NM_0808	Homo sapiens endothelial lectin HL-2 (HL-2), mRNA Homo sapiens protein phasely to the same sapiens protein phasely to the same sapiens protein phasely to the same same same same same same same sam
NM_08087	Homo sapiens protein phanel
NM_08087	
NM 08087	
NM_08087	Homo saniens VIA A 1777
NM_08086	7 Homo saniens suppress (Unc5h4), mRNA
NM_08086	
NM_08086	Homo sapiens relaxin 3 (H3) (RLN3), mRNA  Homo sapiens aplacia
NM_080862	
	Homo sapiens SPRY domain-containing SOCS box-containing 16 (ASB16), mRNA mRNA
NM_080861	Homo garia appro-
	Homo sapiens SPRY domain-containing SOCS box protein SSB-3 (SSB-3),
NM_080860	Home see:
NM_016150	
NM_016127	Homo sapiens ankyrin repeat and SOCS box-containing 2 (ASB2), mRNA  Homo sapiens hypothetical protein MGC8721 (MGC8721)
NM_004170	Homo sapiens hypothetical protein MGC8721 (MGC8721), mRNA Homo sapiens solute carrier family 1 (neuronal)
1 - 1111_004170	Homo sapiens solute carrier family 1 (neuronal/epithelial high affinity glutamate transporter, system Xag), member 1 (SLC1A1), nuclear games
	transporter, system Xag), member 1 (SLC1A1), nuclear gene encoding mitochondrial protein, mRNA
NM_017611	illitochondrial protein make.
NM_025220	110IIIO Sapiens hymothetical
1111_023220	Homo sapiens a disintegrin and metalloproteinase domain 33 (ADAM33),
NM_018548	I III(I)A
NM_080740	riomo saniens doum re-il i li
141/1_080/40	Homo sapiens similar to Ovis aries V observed
ND C CLOSE	(3'OY11.1), mRNA
NM 012163	riomo saniens F-boy and I
NM_012304	Homo sapiens F-box and leucing rich repeat protein 9 (FBXL9), mRNA
NM_012160	Homo sapiens F-box and level in February (FBXL7), mRNA
NM_012159	Homo sapiens F-box and Issued Homo s
VM_012158	Homo sapiens F-box and level repeat protein 3B (FBXL3B), mRNA
VM_012157	Homo sapiens F-hox and love in the protein 3A (FBXL3A) mRNA
VM_024555	Homo sapiens F-box and leucine-rich repeat protein 3A (FBXL3A), mRNA  Homo sapiens F-box and leucine-rich repeat protein 2 (FBXL2), mRNA
	Homo sapiens F-box and leucine-rich repeat protein 2 (FBXL2), mRNA variant 2, mRNA
VM_012162	Homo sapiens E-how and 1
	Homo sapiens F-box and leucine-rich repeat protein 6 (FBXL6), transcript variant 1, mRNA
IM_033535	Homo sanione E 1
_	Homo sapiens F-box and leucine-rich repeat protein 5 (FBXL5), transcript variant 2, mRNA
M_012161	Homo conica Di
2101	Homo sapiens F-box and leucine-rich repeat protein 5 (FBXL5), transcript variant 1, mRNA
M_002278	Wallall I, mKNA (FBXL5), transcript
	Homo sapiens keratin hoir and in a grand
W_U33283	Homo sapiens tumor protein p53 inducible nuclear protein 1 (TP53INP1),
VI 000077	mRNA rotein 1 (TP53INP1),
VI_002211	Homo sapiens keratin bei
M_032994	Homo sapiens Williams Beuren syndrome chromosome region 14 (WBSCR14),

	transcript variant 5, mRNA
NM 032954	Homo sapiens Williams Beuren syndrome chromosome region 14 (WBSCR14),
14141_032934	transcript variant 4, mRNA
ND4_022052	Homo sapiens Williams Beuren syndrome chromosome region 14 (WBSCR14),
NM_032953	transcript variant 3, mRNA
NM 032952	Homo sapiens Williams Beuren syndrome chromosome region 14 (WBSCR14),
14141_032932	transcript variant 2, mRNA
ND4 022051	Homo sapiens Williams Beuren syndrome chromosome region 14 (WBSCR14),
NM_032951	transcript variant 1, mRNA
NG_000008	Homo sapiens genomic cytochrome P450, subfamily IIA (phenobarbital-
140_00000	inducible) (CYP2A) on chromosome 19
NM 030809	Homo sapiens chromosome 12 open reading frame 22 (C12orf22), mRNA
NM 004426	Homo sapiens early development regulator 1 (polyhomeotic 1 homolog) (EDR1),
1NM_004420	mRNA
ND4 020244	Homo sapiens choline phosphotransferase 1 (CHPT1), mRNA
NM_020244 NM_019074	Homo sapiens delta-like 4 (Drosophila) (DLL4), mRNA
	Homo sapiens chromosome X open reading frame 9 (CXorf9), mRNA
NM_018990	Homo sapiens chromosome 21 open reading frame 55 (C21orf55), mRNA
NM_017833	
NM_018255	Homo sapiens elongator protein 2 (ELP2), mRNA
NM_014096	Homo sapiens hypothetical protein DKFZp762A227 (DKFZp762A227), mRNA
NM_014927	Homo sapiens connector enhancer of KSR2 (CNK2), mRNA
NM_012164	Homo sapiens F-box and WD-40 domain protein 2 (FBXW2), mRNA
NM_012247	Homo sapiens selenium donor protein (SPS), mRNA
NM_012165	Homo sapiens F-box and WD-40 domain protein 3 (FBXW3), mRNA
NM_007198	Homo sapiens proline synthetase co-transcribed homolog (bacterial) (PROSC), mRNA
NM_006011	Homo sapiens sialyltransferase 8B (alpha-2, 8-sialytransferase) (SIAT8B),
14M_000011	mRNA
NM 005674	Homo sapiens zinc finger protein 239 (ZNF239), mRNA
NM 001364	Homo sapiens discs, large homolog 2, chapsyn-110 (Drosophila) (DLG2),
14112_001501	mRNA
NM 000646	Homo sapiens amylo-1, 6-glucosidase, 4-alpha-glucanotransferase (glycogen
1 11.1_0000 10	debranching enzyme, glycogen storage disease type III) (AGL), transcript variant
	6, mRNA
NM 000645	Homo sapiens amylo-1, 6-glucosidase, 4-alpha-glucanotransferase (glycogen
_	debranching enzyme, glycogen storage disease type III) (AGL), transcript variant
	5, mRNA
NM 000644	Homo sapiens amylo-1, 6-glucosidase, 4-alpha-glucanotransferase (glycogen
_	debranching enzyme, glycogen storage disease type III) (AGL), transcript variant
ļ	2, mRNA
NM 000643	Homo sapiens amylo-1, 6-glucosidase, 4-alpha-glucanotransferase (glycogen
	debranching enzyme, glycogen storage disease type III) (AGL), transcript variant
	3, mRNA
NM_000642	Homo sapiens amylo-1, 6-glucosidase, 4-alpha-glucanotransferase (glycogen
_	debranching enzyme, glycogen storage disease type III) (AGL), transcript variant
	1, mRNA
NM_000028	Homo sapiens amylo-1, 6-glucosidase, 4-alpha-glucanotransferase (glycogen
-	debranching enzyme, glycogen storage disease type III) (AGL), transcript variant
1	000101111111   01111
	4, mRNA
NM_080831	4, mRNA
NM_080831 NM_080825	

NIM DITTE	PC 1/US03/050
NM_01766	
NM_08074	6 (TRPM6), mRNA
- 1212_00074	
NM_00049	domains) (SRCRB4D), mRNA
	suproms comagen time x alpha 1/0-1
NM_05709	(COL10A1), mRNA  6 Home conjugate to the first term of the first t
	Homo sapiens cytochrome P450 polypeptide 43 (CYP3A43), transcript variant 3
NM_014578	Homo seniene and I
NM_020708	
	Homo sapiens solute carrier family 12, (potassium-chloride transporter) member 5 (SLC12A5), mRNA
NM_016093	Homo saniens ribeassed
NM_057095	
	mRNA polypeptide 43 (CYP3A43), transcript variant 2
NM_022820	Homo saniens cytochrow PASO
	Homo sapiens cytochrome P450 polypeptide 43 (CYP3A43), transcript variant 1
NM_052969	Homo saniens ribosomal and in Your
NM_052970	Homo sapiens ribosomal protein L39-like (RPL39L), mRNA Homo sapiens chromosome 20 cm.
NM_052865	Homo sapiens chromosome 20 open reading frame 60 (C20orf60), mRNA  Homo sapiens chromosome 20 open reading frame 70 (C20orf60), mRNA
NM_021029	Homo sapiens chromosome 20 open reading frame 60 (C20orf60), mRNA  Homo sapiens ribosomal protein I 36a (RPI 36A), PNA
NM_001001	Homo sapiens ribosomal protein L36a (RPL36A), mRNA Homo sapiens ribosomal protein L36a (RPL36A), mRNA
NM_033645	
	Homo sapiens F-box and WD-40 domain protein 1B (FBXW1B), transcript variant 1, mRNA
NM_033644	Homo saniens E have a vivo
	Homo sapiens F-box and WD-40 domain protein 1B (FBXW1B), transcript variant 2, mRNA
NM_012300	Homo saniens E home 1 MD
	Homo sapiens F-box and WD-40 domain protein 1B (FBXW1B), transcript variant 3, mRNA
NM_022760	Homo saniens chromoso 20
NM_014958	Homo sapiens chromosome 20 open reading frame 81 (C20orf81), mRNA  Homo sapiens Rho guanine nucleotide exchange for the CONTRA
	Homo sapiens Rho guanine nucleotide exchange factor (GEF) 15 (ARHGEF15),
NM_021810	Homo saniens cadharin 111 26 (or
NM_030876	Homo sapiens cadherin-like 26 (CDH26), mRNA
	Homo sapiens olfactory receptor, family 5, subfamily V, member 1 (OR5V1),
NM_031232	Homo saniens amyloid hote (A.4)
	Homo sapiens amyloid beta (A4) precursor protein-binding, family A, member 2 binding protein (APBA2BP), transcript variant 2 mPNA
NM_031231	binding protein (APBA2BP), transcript variant 2, mRNA  Homo sapiens amyloid bets (A4)
NM_032554	binding protein (APBA2BP), transcript variant 1, mRNA
NM_006462	Homo sapiens G protein-coupled receptor 81 (GPR81), mRNA  Homo sapiens chromosome 20 cner at 15 (GPR81), mRNA
	variant 1, mRNA
VM_031229	Homo sapiens chromosome 20 open reading frame 18 (C20orf18), transcript variant 2, mRNA
	variant 2, mRNA
VM_031228	Homo sapiens chromosome 20
	Homo sapiens chromosome 20 open reading frame 18 (C20orf18), transcript variant 3, mRNA
JM_031227	Homo sapiens chromosome 20 organization
	Homo sapiens chromosome 20 open reading frame 18 (C20orf18), transcript variant 4, mRNA
M_031424	Homo sapiens chromosome 20 annu i
M_000518	Homo sapiens chromosome 20 open reading frame 55 (C20orf55), mRNA Homo sapiens hemoglobin, beta (HBB), mRNA
M_030959	Homo sapiens olfactory recenter for its the sapiens of the sapiens
2.12_030333	Homo sapiens olfactory receptor, family 12, subfamily D, member 3 (OR12D3),
M_018661	Homo sapiens defensin hete 2 (DEPDE)
M_018661	Homo sapiens defensin, beta 3 (DEFB3), mRNA Homo sapiens DNA cross-link repair 1C (PSO2 homolog, S. cerevisiae)

	(DCLRE1C), mRNA
NM 022099	Homo sapiens chromosome 20 open reading frame 51 (C20orf51), mRNA
NM 000668	Homo sapiens alcohol dehydrogenase IB (class I), beta polypeptide (ADH1B),
	mRNA
NM 021943	Homo sapiens testis expressed sequence 27 (TEX27), mRNA
NM_021640	Homo sapiens chromosome 12 open reading frame 10 (C12orf10), mRNA
NM_021215	Homo sapiens chromosome 20 open reading frame 77 (C20orf77), mRNA
NM_012141	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 26 (DDX26),
	mRNA
NM_021225	Homo sapiens proline-rich 1 (PROL1), mRNA
NM_006508	Homo sapiens regenerating islet-derived-like, pancreatic stone protein-like,
	pancreatic thread protein-like (rat) (REGL), mRNA
NM_020356	Homo sapiens chromosome 20 open reading frame 32 (C20orf32), mRNA
NM_020369	Homo sapiens fascin homolog 3, actin-bundling protein, testicular
	(Strongylocentrotus purpuratus) (FSCN3), mRNA
NM_020145	Homo sapiens SH3-domain GRB2-like endophilin B2 (SH3GLB2), mRNA
NM_020125	Homo sapiens BCM-like membrane protein precursor (BLAME), mRNA
NM_019025	Homo sapiens chromosome 20 open reading frame 16 (C20orf16), mRNA
NM_018679	Homo sapiens t-complex 11 (mouse) (TCP11), mRNA
NM_017589	Homo sapiens B-cell translocation gene 4 (BTG4), mRNA
NM_018692	Homo sapiens chromosome 20 open reading frame 17 (C20orf17), mRNA
NM_018697	Homo sapiens LanC lantibiotic synthetase component C-like 2 (bacterial)
ND 6 019677	(LANCL2), mRNA
NM_018677	Homo sapiens acetyl-Coenzyme A synthetase 2 (ADP forming) (ACAS2), mRNA
NM_018431	Homo sapiens chromosome 20 open reading frame 180 (C20orf180), mRNA
NM_018725	Homo sapiens interleukin 17B receptor (IL17BR), mRNA
NM 018474	Homo sapiens chromosome 20 open reading frame 19 (C20orf19), mRNA
NM 018478	Homo sapiens chromosome 20 open reading frame 35 (C20orf35), mRNA
NM 017896	Homo sapiens chromosome 20 open reading frame 11 (C20orf11), mRNA
NM 017874	Homo sapiens chromosome 20 open reading frame 27 (C20orf27), mRNA
NM 017859	Homo sapiens uridine kinase-like 1 (URKL1), mRNA
NM 017798	Homo sapiens chromosome 20 open reading frame 21 (C20orf21), mRNA
NM_017789	Homo sapiens sema domain, immunoglobulin domain (Ig), transmembrane
	domain (TM) and short cytoplasmic domain, (semaphorin) 4C (SEMA4C),
	mRNA
NM_017714	Homo sapiens chromosome 20 open reading frame 13 (C20orf13), mRNA
NM_017671	Homo sapiens chromosome 20 open reading frame 42 (C20orf42), mRNA
NM_018384	Homo sapiens immune associated nucleotide 4 like 1 (mouse) (IAN4L1), mRNA
NM_018354	Homo sapiens chromosome 20 open reading frame 46 (C20orf46), mRNA
NM_018347	Homo sapiens chromosome 20 open reading frame 29 (C20orf29), mRNA
NM_018327	Homo sapiens chromosome 20 open reading frame 38 (C20orf38), mRNA
NM_018282	Homo sapiens paraspeckle protein 1 (PSP1), mRNA
NM_018270	Homo sapiens chromosome 20 open reading frame 20 (C20orf20), mRNA
NM_018257	Homo sapiens chromosome 20 open reading frame 36 (C20orf36), mRNA
NM_018197	Homo sapiens zinc finger protein 64 homolog (mouse) (ZFP64), mRNA
NM_018010	Homo sapiens estrogen-related receptor beta like 1 (ESRRBL1), mRNA
NM_017446	Homo sapiens mitochondrial ribosomal protein L39 (MRPL39), mRNA
NM_017429	Homo sapiens beta-carotene 15, 15'-dioxygenase (BCDO), mRNA
NM_016082	Homo sapiens chromosome 20 open reading frame 34 (C20orf34), mRNA
NM_016610	Homo sapiens toll-like receptor 8 (TLR8), mRNA
NM_016009	Homo sapiens SH3-domain GRB2-like endophilin B1 (SH3GLB1), mRNA

ND C OF CO.	PC1/US03/050
NM_01640	
NM_01640	
NM_016319	Homo sapiens COP9 constitutive photomorphogenic homolog subunit 7A  (Arabidopsis) (COPS7A) mPNA
NTM 01500	(Arabidopsis) (COPS7A), mRNA
NM_015985	Flomo sapiens angionojetin 4 (ANCETA) PAR
NM_015834	Tionio sapiciis adenosine deaminase Data
	(ADARB1), transcript variant DRADA2c, mRNA  Homo sanians administration of the sanians and the sanians administration of the sanians and the sanians administration of the sanians adminis
NM_015833	Homo sapiens adenosine deaminaco DNA
	Homo sapiens adenosine deaminase, RNA-specific, B1 (RED1 homolog rat)  (ADARB1), transcript variant DRABA2b, mRNA
NM_014036	Homo sapiens BCM-like membrana mada
NM_014012	Homo sapiens RAS (RAD and GEM)-like GTP-binding (REM), mRNA Homo sapiens synaptosomal-associated
NM_014841	Homo sapiens synantosomal associated
	(SNAP91), mRNA (SNAP91), mRNA
NM_014795	Homo sapiens zinc finger homosham 11 (GRAVIII)
NM_015313	Homo sapiens Rho guanine nucleotide exchange factor (GEF) 12 (ARHGEF12)
	mRNA mRNA
NM_014784	Homo sapiens Rho guanine nucleotide exchange factor (GEF) 11 (ARHGEF11)
	mRNA mRNA
NM_014862	Homo saniens and hydrogod
	Homo sapiens aryl-hydrocarbon receptor nuclear translocator 2 (ARNT2), mRNA
NM_014054	Homo saniens chromosom 20
NM_015629	Homo sapiens chromosome 20 open reading frame 40 (C20orf40), mRNA Homo sapiens PRP31 pre-mRNA processing for the 21th
	Homo sapiens PRP31 pre-mRNA processing factor 31 homolog (yeast)  (PRPF31), mRNA
NM_015417	Homo saniene abronous as
NM_014625	Homo sapiens chromosome 20 open reading frame 28 (C20orf28), mRNA  Homo sapiens nephrosis 2, idiopathic steroid and the control of the contro
_	Homo sapiens nephrosis 2, idiopathic, steroid-resistant (podocin) (NPHS2), mRNA
VM_014592	Homo saniens Ky change 1
VM_014140	Homo sapiens Kv channel interacting protein 1 (KCNIP1), mRNA  Homo sapiens SWI/SNE related many interacting protein 1 (KCNIP1), mRNA
-	Homo sapiens SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a-like 1 (SMARCALL) mPNA
VM_013442	chromatin, subfamily a-like 1 (SMARCAL1), mRNA  Homo saniens stometin (EPDZO), iii
VM_013248	
VM 013316	
JM_013348	
	Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 14 (KCNJ14), mRNA
M_013279	Home carried 1
M_012418	Homo sapiens chromosome 11 open reading frame 9 (C11orf9), mRNA  Homo sapiens fascin homolog 2 patie have 11
	Homo sapiens fascin homolog 2, actin-bundling protein, retinal
M_012201	(Strongylocentrotus purpuratus) (FSCN2), mRNA
M 000519	The state of the s
M_006999	The supreme delta (HDD) - Data
M_006719	TIOMO Sapiciis Dolymerase (INTA 1: 1)
	Homo sapiens actin binding LIM protein (ABLIM), transcript variant ABLIM-m,
M_002313	TIRINA TIL
002515	Homo sapiens actin binding LIM protein (ABLIM), transcript variant ABLIM-l, mRNA
<u>1_007238</u>	TICLUS - CALLES THE VARIANT ABLIM-I,
_	Homo sapiens peroxisomal membrane protein 4 (24kD) (PXMP4), mRNA Homo sapiens nischarin (NISCH), mRNA
4_006720	Homo sapiens nischarin (NISCH), mRNA
_000720	Tromo sapiens actin binding I IM protein (A DI D.C.
1_007026	mRNA protein (ABLIM), transcript variant ABLIM-s,
1_00/026	Homo sapiens dual specificity at a discontinuo
	Archidensia (Consultation Photomolog Submit 5
1_006614 I	(Arabidopsis) (COPS5), mRNA  Homo sapiens cell adhesion molecule with homology to L1CAM (close homolog

	of 1) (CIII 1) mDNIA
ND 4 00 6 4 1 0	of L1) (CHL1), mRNA
NM_006410	Homo sapiens HIV-1 Tat interactive protein 2, 30 kD (HTATIP2), mRNA
NM_006432	Homo sapiens Niemann-Pick disease, type C2 (NPC2), mRNA Homo sapiens golgi transport complex 1 (90 kD subunit) (GOLTC1), mRNA
NM_006348	Homo sapiens goigi transport complex 1 (90 kD subulit) (GOLTCT), filkANA
NM_006408	Homo sapiens anterior gradient 2 homolog (Xenepus laevis) (AGR2), mRNA
NM_006106	Homo sapiens Yes-associated protein 1, 65 kD (YAP1), mRNA
NM_006096	Homo sapiens N-myc downstream regulated gene 1 (NDRG1), mRNA
NM_006071	Homo sapiens polycystic kidney disease (polycystin) and REJ (sperm receptor for egg jelly homolog, sea urchin)-like (PKDREJ), mRNA
NM 006092	Homo sapiens caspase recruitment domain family, member 4 (CARD4), mRNA
NM 005748	Homo sapiens YY1 associated factor 2 (YAF2), mRNA
NM 005715	Homo sapiens uronyl-2-sulfotransferase (UST), mRNA
NM 005622	Homo sapiens SA hypertension-associated homolog (rat) (SAH), mRNA
NM_005733	Homo sapiens RAB6 interacting, kinesin-like (rabkinesin6) (RAB6KIFL), mRNA
NM_005668	Homo sapiens sialyltransferase 8D (alpha-2, 8-polysialytransferase) (SIAT8D), mRNA
NM 005606	Homo sapiens legumain (LGMN), mRNA
NM 004649	Homo sapiens chromosome 21 open reading frame 33 (C21orf33), mRNA
NM 005469	Homo sapiens peroxisomal acyl-CoA thioesterase (PTE1), mRNA
NM 005180	Homo sapiens B lymphoma Mo-MLV insertion region (mouse) (BMI1), mRNA
NM 005108	Homo sapiens xylulokinase homolog (H. influenzae) (XYLB), mRNA
NM_004610	Homo sapiens t-complex 10 (mouse) (TCP10), mRNA
NM_004579	Homo sapiens mitogen-activated protein kinase kinase kinase kinase 2 (MAP4K2), mRNA
NM_004086	Homo sapiens coagulation factor C homolog, cochlin (Limulus polyphemus)
	(COCH), mRNA
NM_004273	Homo sapiens carbohydrate (chondroitin 6) sulfotransferase 3 (CHST3), mRNA
NM_004902	Homo sapiens RNA-binding region (RNP1, RRM) containing 2 (RNPC2), mRNA
NM 004353	Homo sapiens serine (or cysteine) proteinase inhibitor, clade H (heat shock
	protein 47), member 1, (collagen binding protein 1) (SERPINH1), mRNA
NM_004317	Homo sapiens arsA arsenite transporter, ATP-binding, homolog 1 (bacterial) (ASNA1), mRNA
NM_001247	Homo sapiens ectonucleoside triphosphate diphosphohydrolase 6 (putative function) (ENTPD6), mRNA
NM_003831	Homo sapiens sudD suppressor of bimD6 homolog (A. nidulans) (SUDD), mRNA
NM 003143	Homo sapiens single-stranded DNA binding protein (SSBP1), mRNA
NM 003098	Homo sapiens syntrophin, alpha 1 (dystrophin-associated protein A1, 59kD,
	acidic component) (SNTA1), mRNA
NM_003034	Homo sapiens sialyltransferase 8A (alpha-N-acetylneuraminate/alpha-2,8-sialytransferase, GD3 synthase) (SIAT8A), mRNA
NM_003028	Homo sapiens SHB (Src homology 2 domain-containing) adaptor protein B (SHB), mRNA
NM 003579	Homo sapiens RAD54-like (S. cerevisiae) (RAD54L), mRNA
NM_002669	Homo sapiens pleiotropic regulator 1 (PRL1homolog, Arabidopsis) (PLRG1), mRNA
NM_000139	Homo sapiens membrane-spanning 4-domains, subfamily A, member 1 (MS4A2), mRNA
NM_003836	Homo sapiens delta-like 1 homolog (Drosophila) (DLK1), mRNA
NM 003653	Homo sapiens COP9 constitutive photomorphogenic homolog subunit 3
14141 002022	1 Monte supreme CO1 > Constitutive photomorphogenic nomotog subulit 3

	(Arabidonsis) (COPS2)
NM_000083	(Arabidopsis) (COPS3), mRNA
	dominant) (CLCN1) - Party
NM_000691	dominant) (CLCN1), mRNA
_	Homo sapiens aldehyde dehydrogenase 3 family, memberA1 (ALDH3A1), mRNA
NM_001112	Home social distribution,
NM_004370	(ADARB1), transcript variant DRADA2a, mRNA  Homo saniens collager to ANA
1370	Homo sapiens collagen, type XII, alpha 1 (COL12A1), transcript variant long, mRNA
NM_080645	Home variant long,
	Homo sapiens collagen, type XII, alpha 1 (COL12A1), transcript variant short,
NM_080681	United the same of
	Homo sapiens collagen, type XI, alpha 2 (COL11A2), transcript variant 2,
NM_080680	Home and the transcript variant 2,
- 1212_000000	Homo sapiens collagen, type XI, alpha 2 (COL11A2), transcript variant 1,
NM_080679	The transcript variant I,
- 44_0000/9	Homo sapiens collagen, type XI, alpha 2 (COL11A2), transcript variant 3,
NM_003593	mkina 1 = (00D11A2), uanscript variant 3,
NM_000638	Homo sapiens winged-helix nude (WHN), mRNA
14141_000038	
NM_080682	Homo sapiens vitronectin (serum spreading factor, somatomedin B, complement S-protein) (VTN), mRNA
T41AT_090095	Homo sapiens vascular cell adhesion malantal discon-
NM_001078	mRNA addressor molecule I (VCAM1), transcript variant 2,
T4141_0010\8	Homo sapiens vascular cell adhesion molecule 1 (VCAM1), transcript variant 1,
NM 000115	mknA molecule 1 (VCAIVII), transcript variant 1,
NM_006115	Homo sapiens preferentially over the same same same same same same same sam
NM_000175	Homo sapiens glucose phosphate isomerase (GPI), mRNA  Homo sapiens EnhAS (EDHAS), DNA
NM_020526	Homo sapiens EphAs (FPHAs) - PNIA
NM_002109	Homo sapiens histidyl-tRNA gymth-tagget
NM_012208	Homo sapiens histidyl-tRNA synthetase-like (HARSL), mRNA Homo sapiens T-box 6 (TRY6)
NM_004608	Homo sapiens T-box 6 (TBX6), transcript variant 1, mRNA  Homo sapiens T-box 6 (TBX6), transcript variant 1, mRNA
VM_080758	Homo sapiens T-box 6 (TBX6), transcript variant 1, mRNA Homo sapiens T-box 5 (TBX5), transcript variant 2, mRNA
VM_080718	Homo sapiens T-box 5 (TBX5), transcript variant 2, mRNA Homo sapiens T-box 5 (TBX5), transcript variant 2, mRNA
VM_080717	Homo sapiens T-boy 5 (TDV5), transcript variant 2, mRNA
M_000192	Homo sapiens T-boy 5 (TDV5), transcript variant 3, mRNA
VM_080832	Homo sapiens T-box 5 (TBX5), transcript variant 3, mRNA Homo sapiens poly(A) binding protein
IM_080824	Homo sapiens poly(A) binding protein, cytoplasmic 5 (PABPC5), mRNA Homo sapiens chromosome 20 open reading frame 106 (CR)
IM_080822	Homo sapiens chromosome 20 open reading frame 106 (C20orf106), mRNA Homo sapiens candidate tumor suppressor OVCA 2 (OVCA 2)
M_080821	Homo sapiens candidate tumor suppressor OVCA2 (OVCA2), mRNA Homo sapiens chromosome 20 open real in School (OVCA2), mRNA
M_080820	Homo sapiens chromosome 20 open reading frame 108 (C20orf108), mRNA Homo sapiens chromosome 20 open reading frame \$\$ (C20orf108), mRNA
M_080818	Homo sapiens chromosome 20 open reading frame 108 (C20orf108), mRNA Homo sapiens G protein-coupled recentor 80 (CPR 88), mRNA
M_080817	Homo sapiens G protein-coupled receptor 80 (GPR80), mRNA Homo sapiens G protein-coupled receptor 80 (GPR80), mRNA
M_080794	Homo sapiens G protein-coupled receptor 80 (GPR80), mRNA Homo sapiens mitochondrial ribesee the same same same same same same same sam
M_020973   I	Homo sapiens mitochondrial ribosomal protein L39 (MRPL39), mRNA Homo sapiens cytosolic beta-glucosidasa (GLHG).
M 054112 H	Homo sapiens cytosolic beta-glucosidase (GLUC), mRNA  Homo sapiens chromosoma 20
$M_014145$ F	Homo sapiens chromosome 20 open reading frame 63 (C20orf63), mRNA Homo sapiens chromosome 20 open reading frame 167 (C20orf167), mRNA
M 033409 H	Homo sapiens chromosome 20 open reading frame 167 (C20orf167), mRNA Homo sapiens chromosome 20 open reading frame 30 (C20orf30), mRNA
1 033409 F	Iomo sapiens chromosome 20 oper realing frame 30 (C20orf30), mRNA
VI_032013 H	Iomo sapiens NDRG family more leading frame 34 (C20orf54), mRNA
1 032109 H	omo sapiens orthopedia homela (C) (NDKOS), MRNA
(1)	omo sapiens membrane-spanning 4-domains, subfamily A, member 4  MS4A4A), mRNA
1_022910 H	omo sapiens NDRG family member 4 (NDRG4), mRNA

3), mRNA 9), mRNA 149), mRNA abunit 7B Q1), mRNA
9), mRNA 149), mRNA 15 15 16 17 18 18 18 18 18 18 18 18 18 18 18 18 18
ubunit 7B Q1), mRNA
ubunit 7B Q1), mRNA
ubunit 7B Q1), mRNA
Q1), mRNA
LDH8A1),
9), mRNA
oorter-like,
,
MLL3), mRNA
DMTF1),
,,,
og 1 (avian)
nsporter),
, , , , , , , , , , , , , , , , , , ,
A
(Allgrove, triple-
ember 2 (Fc
(MS4A1), mRNA
ALDH6A1),
ase, spastic
, o, op
IRC6), mRNA
1), mRNA
ne homolog F
s) (NUDC),
J, (21020),
(DNAJB1),
(~~11.10.2.1),
ed protein, 230
protein, 250
hthamide
f143), mRNA
,,
de 27 (DDX27),
9), mRNA
2), mRNA
35), mRNA
A

\*.

1	
NM_08074	
NM_080739	TIVITO SADICIIS CHTOMOCOme 20
NM_033550	Homo sapiens chromosome 20 open reading frame 141 (C20orf141), mRNA
NM_080732	Homo sapiens chromosome 20 open reading frame 141 (C20orf141), mRNA  Homo sapiens egl nine homolog 2 (C. elegens) (ECLNO)
277	mRNA (C. clegalis) (EGLN2), transcript variant 3
NM_053046	Homo sapiens egl nine homolog 2 (C. 1
	Homo sapiens egl nine homolog 2 (C. elegans) (EGLN2), transcript variant 1,
NM_025106	Homo sapiens SPRY domain acertain GO Con-
	Homo sapiens SPRY domain-containing SOCS box protein SSB-1 (FLJ22393),
NM_030760	Homo sapiens endothelial differentiation, sphingolipid G-protein-coupled receptor, 8 (EDG8), mRNA
	receptor, 8 (EDG8), mRNA
NM_016069	Homo sapjens mitochondria and it is
	macrophage colony-stimulating factors involved in granulocyte-
	gene encoding mitochondriel and signal transduction (Magmas), nuclear
NM_021205	Homo sapiens Wnt-1 responsive Cd. 404
NM_032495	Homo sapiens hypothetical protein SMAP31 (SMAP31), mRNA Homo sapiens interleukin-1 HV2 (R. HNV2)
NM_032556	Homo sapiens interleukin 1 LIV2 (M. 11VIV) (SMAP31), mRNA
NM_014331	
	system) member 11 (SLC7A11), mRNA
NM_017564	Homo sapiens stabilin-2 (STAB2), mRNA Homo sapiens his a Galiana S
NM_020924	1 - 20110 Sapicils Dioref (bioref) - Data
NM_015356	Homo sapiens scribble (SCRIB), mRNA
NM_030648	1 2101110 Sapiells SEI domain contain
NM_018488	Homo sapiens SET domain-containing protein 7 (SET7), mRNA  Homo sapiens T-box 4 (TBX4), mRNA
NM_016470	Homo sapiens chromosome 20 map 20q13.11
NM_080722	Homo sapiens a disintegrin 111
	Homo sapiens a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif. 14 (ADAMTS14) ar DNA
NM_080676	Homo sapiens chromosome 20
NM_080674	Homo sapiens chromosome 20 open reading frame 133 (C20orf133), mRNA Homo sapiens chromosome 20 open reading frame 86 (C20orf86), mRNA
NM_080621	Homo sapiens chromosome 20 - February Hame 86 (C20ort86), mRNA
NM_080608	Homo sapiens chromosome 20 open reading frame 136 (C20orf136), mRNA Homo sapiens hypothetical protein MGC4473 (MGC4473)
NM_080719	Homo sapiens hypothetical and 100 (C20orf165), mRNA
NM_003495	Homo sapiens hypothetical protein MGC4473 (MGC4473), mRNA Homo sapiens H4 histone family, marsh as MGC4473), mRNA
NM_020633	Homo sapiens V1R-like 1 (V1RL1), mRNA  Homo sapiens W1R-like 1 (V1RL1), mRNA
	Homo sapjens vacuolar modeli, mikiva
NM_080631	Homo sapiens vacuolar protein sorting 45A (yeast) (VPS45A), mRNA Homo sapiens vacuolar protein sorting 41 (yeast) (VPS45A), mRNA
j	Homo sapiens vacuolar protein sorting 45A (yeast) (VPS45A), mRNA mRNA
NM_014396	Homo saniens vacuolar made
	Homo sapiens vacuolar protein sorting 41 (yeast) (VPS41), transcript variant 1,
VM_018668	Homo saniens vacualer mode:
VM_022916	Homo sapiens vacuolar protein sorting 33B (yeast) (VPS33B), mRNA
M_003610 ]	Homo sapiens vacuolar protein sorting 33B (yeast) (VPS33B), mRNA Homo sapiens RAE1 RNA export 1 homolog (S. nombo) (VPS33A), mRNA
1V1_014061   I	Homo sapiens RAE1 RNA export 1 homolog (S. pombe) (RAE1), mRNA Homo sapiens APR-1 protein (MAGEH1), mRNA
IM_001927 H	Homo sapiens APR-1 protein (MAGEH1), mRNA  Homo sapiens desmin (DPS)
	- STATE SUPPLIES (LESITION / ) INDIVI
	Homo sapiens histone family member (H2B/S), mRNA
M_001866 H	ncoding mitochnodrial protein, mRNA
	onio sapiens cylochrome a conid
M_004718 H	ncoding mitochondrial protein, mRNA omo sapiens cytochrome c oxidase subunit VIIb (COX7B), nuclear gene COX7A2L), nuclear gene encoding mitochondrial polypeptide 2 like
	Onio Sapiciis Cyrochromo a - 11
	COX7A2L), nuclear gene encoding mitochondrial protein, mRNA

NM_001865	Homo sapiens cytochrome c oxidase subunit VIIa polypeptide 2 (liver) (COX7A2), nuclear gene encoding mitochondrial protein, mRNA
	(COA/AZ), nuclear gene encounting into condition protein, including
NM_001864	Homo sapiens cytochrome c oxidase subunit VIIa polypeptide 1 (muscle) (COX7A1), nuclear gene encoding mitochondrial protein, mRNA
NM_006438	Homo sapiens collectin sub-family member 10 (C-type lectin) (COLEC10),
14141_000428	mRNA
NM_080544	Homo sapiens collagen-like tail subunit (single strand of homotrimer) of
_	asymmetric acetylcholinesterase (COLQ), transcript variant VIII, mRNA
NM_080543	Homo sapiens collagen-like tail subunit (single strand of homotrimer) of
111120000 10	asymmetric acetylcholinesterase (COLQ), transcript variant VII, mRNA
NM_080542	Homo sapiens collagen-like tail subunit (single strand of homotrimer) of
14141_000342	asymmetric acetylcholinesterase (COLQ), transcript variant VI, mRNA
ND 4 000541	Homo sapiens collagen-like tail subunit (single strand of homotrimer) of
NM_080541	
77.5.000540	asymmetric acetylcholinesterase (COLQ), transcript variant V, mRNA
NM_080540	Homo sapiens collagen-like tail subunit (single strand of homotrimer) of
	asymmetric acetylcholinesterase (COLQ), transcript variant IV, mRNA
NM_080539	Homo sapiens collagen-like tail subunit (single strand of homotrimer) of
	asymmetric acetylcholinesterase (COLQ), transcript variant III, mRNA
NM_080538	Homo sapiens collagen-like tail subunit (single strand of homotrimer) of
	asymmetric acetylcholinesterase (COLQ), transcript variant II, mRNA
NM_005677	Homo sapiens collagen-like tail subunit (single strand of homotrimer) of
_	asymmetric acetylcholinesterase (COLQ), transcript variant I, mRNA
NM_080592	Homo sapiens apoptosis related protein APR-3 (APR-3), transcript variant 2,
_	mRNA
NM_016085	Homo sapiens apoptosis related protein APR-3 (APR-3), transcript variant 1,
	mRNA
NM_014318	Homo sapiens apoptosis related protein (APR-2), mRNA
NM_001745	Homo sapiens calcium modulating ligand (CAMLG), mRNA
NM_004341	Homo sapiens carbamoyl-phosphate synthetase 2, aspartate transcarbamylase, and dihydroorotase (CAD), nuclear gene encoding mitochondrial protein,
	mRNA
NM_032493	Homo sapiens adaptor-related protein complex 1, mu 1 subunit (AP1M1),
INIVI_032493	mRNA
NM_001128	Homo sapiens adaptor-related protein complex 1, gamma 1 subunit (AP1G1),
_	mRNA
NM 080545	Homo sapiens adaptor-related protein complex 1, gamma 2 subunit (AP1G2),
1111_0000 10	transcript variant 2, mRNA
NM 003917	Homo sapiens adaptor-related protein complex 1, gamma 2 subunit (AP1G2),
14141_003517	transcript variant 1, mRNA
NM 080549	Homo sapiens protein tyrosine phosphatase, non-receptor type 6 (PTPN6),
14141_000243	transcript variant 3, mRNA
NIM 000540	Homo sapiens protein tyrosine phosphatase, non-receptor type 6 (PTPN6),
NM_080548	
NR 6 000001	transcript variant 2, mRNA
NM_002831	Homo sapiens protein tyrosine phosphatase, non-receptor type 6 (PTPN6),
776 00000	transcript variant 1, mRNA
NM_002830	Homo sapiens protein tyrosine phosphatase, non-receptor type 4
	(megakaryocyte) (PTPN4), mRNA
NM_002829	Homo sapiens protein tyrosine phosphatase, non-receptor type 3 (PTPN3),
	mRNA
NM_080423	Homo sapiens protein tyrosine phosphatase, non-receptor type 2 (PTPN2),
	transcript variant 3, mRNA
NM 080422	Homo sapiens protein tyrosine phosphatase, non-receptor type 2 (PTPN2),
NM_080422	Homo sapiens protein tyrosine phosphatase, non-receptor type 2 (PIPN2),

NIM COOCCE	transcript variant 2, mRNA
NM_00282	Homo sapiens protein tyrosine phosphotose man
377.5	transcript variant 1, mRNA
NM_00282	Homo sapiens protein tyrosine phosphatase, non-receptor type 1 (PTPN1), mRNA
277	mRNA phosphatase, non-receptor type 1 (PTPN1),
NM_01424	The state of the property of the state of th
1	Homo sapiens protein tyrosine phosphatase-like (proline instead of catalytic arginine), member a (PTPLA), mRNA
NM_003479	Homo sapiens protein tyrosine phosphotose by RI
350	Homo sapiens protein tyrosine phosphatase type IVA, member 2 (PTP4A2), transcript variant 1, mRNA
NM_080392	Homo sapiens protein tyrosine phosphotoca to This
377	transcript variant 3, mRNA
NM_080391	Homo sapiens protein tyrosine phosphotogo + W. R.
	Homo sapiens protein tyrosine phosphatase type IVA, member 2 (PTP4A2), transcript variant 2, mRNA
NM_080591	Homo sapiens prostaglandin endongeratil
	synthase and cyclooxygenase) (PTGS1), transcript variant 2, mRNA
NM_000962	Homo sapiens prostaglandin endengani il
	synthase and cyclooxygenase) (PTCC1)
NM_004058	Homo sapiens calcyphosine (CAPS) is transcript variant 1, mRNA
NM_080590	Homo sapiens calcyphosine (CAPS), transcript variant 1, mRNA
NM_006380	
	2 (APPBP2), mRNA
NM_003905	Homo sapiens amyloid beto present
_	Homo sapiens amyloid beta precursor protein binding protein 1, 59kD (APPBP1), mRNA
NM_005783	Homo sapiens ATP binding protein associated with cell differentiation (APACD), mRNA
	(APACD), mRNA
NM_080600	Homo sapiens myelin essecit 1 1
	Homo sapiens myelin associated glycoprotein (MAG), transcript variant 2, mRNA
NM_002361	Homo sapiens myelin eggarid 1
	Homo sapiens myelin associated glycoprotein (MAG), transcript variant 1,
NM_005994	Homo sapiens T-box 2 (TBX2), mRNA
NM_080647	Homo sapiens T-box 1 (TDV1)
NM_080646	Homo sapiens T-box 1 (TBX1), transcript variant C, mRNA  Homo sapiens T-box 1 (TBX1)
VM_080675	
VM_080617	
VM_080611	Homo sapiens dual specificite et al. (CBLNL1), mRNA
M_080610	
M_080602	Homo sapiens cystatin 9-like (mouse) (CST9L), mRNA  Homo sapiens actin related
_	Homo sapiens actin related protein 2/3 complex, subunit 3B (21 kD) (ARPC3B), mRNA
IG_000011	Homo saniens comenia
-	Homo sapiens genomic cytochrome P450, subfamily IIA (phenobarbital-inducible) (CYP2A.3@) on chromosome 10
M_016649	inducible) (CYP2A.3@) on chromosome 19
M_080597	Homo sapiens chromosome 20 open reading frame 6 (C20orf6), mRNA  Homo sapiens oxysterol binding protein libration (C20orf6), mRNA
M_080605	Homo sapiens oxysterol binding protein-like 1A (OSBPL1A), mRNA Homo sapiens UDP-Gal: betaGloNA a beta id a (OSBPL1A), mRNA
	Homo sapiens UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 6 (B3GALT6), mRNA
M_058169	Homo saniana loca (1)
	Homo sapiens loss of heterozygosity, 12, chromosomal region 1 (LOH12CR1), mRNA
M_058164	Homo sanieno elferia di alla d
	Homo sapiens olfactomedin 2 (OLFM2), mRNA
	The support of the su
	The baptells (IX VSIPro) binding mantain 1:1
G_000013	Homo sapiens chorionic gonadotropin, beta polypeptide 7 (CGB7), mRNA Homo sapiens genomic MHC class III complementation
	Homo sapiens genomic MHC class III complement gene cluster (MCGC@) on
	An Uniosome 6

	5 O LOO A 6 1
NM_020967	Homo sapiens nuclear receptor coactivator 5 (NCOA5), mRNA
NM_033044	Homo sapiens microtubule-actin crosslinking factor 1 (MACF1), transcript
	variant 3, mRNA
NM_033024	Homo sapiens microtubule-actin crosslinking factor 1 (MACF1), transcript
·	variant 2, mRNA
NG_000017	Homo sapiens genomic protocadherin beta cluster (PCDHB@) on chromosome 5
NM 015864	Homo sapiens chromosome 6 open reading frame 32 (C6orf32), mRNA
NM 032188	Homo sapiens histone acetyltransferase MYST1 (MYST1), mRNA
NM 030776	Homo sapiens chromosome 20 open reading frame 183 (C20orf183), mRNA
NM 024918	Homo sapiens chromosome 20 open reading frame 172 (C20orf172), mRNA
NM 024812	Homo sapiens brain and acute leukemia, cytoplasmic (BAALC), mRNA
NM 024777	Homo sapiens chromosome 20 open reading frame 124 (C20orf124), mRNA
NM 024758	Homo sapiens agmatinase (FLJ23384), mRNA
NM 024641	Homo sapiens mandaselin (FLJ12838), mRNA
NM 024331	Homo sapiens chromosome 20 open reading frame 121 (C20orf121), mRNA
NM 024301	Homo sapiens fukutin-related protein (FKRP), mRNA
NM 005763	Homo sapiens aminoadipate-semialdehyde synthase (AASS), mRNA
NM 023935	Homo sapiens chromosome 20 open reading frame 116 (C20orf116), mRNA
NM 021993	Homo sapiens FUS interacting protein (serine-arginine rich) 2 (FUSIP2), mRNA
NM_014555	Homo sapiens transient receptor potential cation channel, subfamily M, member
1411_014555	
ND4 000527	5 (TRPM5), mRNA
NM_000537	Homo sapiens renin (REN), mRNA Homo sapiens Crn, crooked neck-like 1 (Drosophila) (CRNKL1), mRNA
NM_016652	
NM_021245	Homo sapiens myozenin 1 (MYOZ1), mRNA
NM_001967	Homo sapiens eukaryotic translation initiation factor 4A, isoform 2 (EIF4A2), mRNA
NM_018649	Homo sapiens H2A histone family, member Y2 (H2AFY2), mRNA
NM_015148	Homo sapiens PAS domain containing serine/threonine kinase (PASK), mRNA
NM_017902	Homo sapiens hypoxia-inducible factor 1, alpha subunit inhibitor (HIF1AN), mRNA
NM_018285	Homo sapiens chromosome 15 open reading frame 12 (C15orf12), nuclear gene
1411_010200	encoding mitochondrial protein, mRNA
NM 018267	Homo sapiens H2A histone family, member J (H2AFJ), mRNA
NM_017555	Homo sapiens egl nine homolog 2 (C. elegans) (EGLN2), transcript variant 2, mRNA
NM 016143	Homo sapiens likely ortholog of rat p47 (p47), mRNA
NM 015993	Homo sapiens plasmolipin (PMLP), mRNA
NM_014938	Homo sapiens Mlx interactor (MONDOA), mRNA  Homo sapiens likely ortholog of mouse ubiquitin conjugating enzyme 7
NM_014948	interacting protein 5 (UBCE7IP5), mRNA
NM_014016	Homo sapiens SAC1 suppressor of actin mutations 1-like (yeast) (SACM1L), mRNA
NM 015156	Homo sapiens REST corepressor (RCOR), mRNA
NM 013337	Homo sapiens translocase of inner mitochondrial membrane 22 homolog (yeast)
	(TIMM22), mRNA
NM 013233	Homo sapiens serine threonine kinase 39 (STE20/SPS1 homolog, yeast)
	(STK39), mRNA
NM 006595	Homo sapiens apoptosis inhibitor 5 (API5), mRNA
NM 006402	Homo sapiens hepatitis B virus x interacting protein (HBXIP), mRNA
NM 006351	Homo sapiens translocase of inner mitochondrial membrane 44 homolog (yeast)
	(TIMM44), mRNA
NM_006327	Homo sapiens translocase of inner mitochondrial membrane 23 homolog (yeast)

Homo sapiens translocase of inner mitochondrial membrane 17 homolog A (yeast) (TIMM17A), mRNA		(TIMM23), mRNA
(yeast) (TMM17A), mRNA  NM_006420  Homo sapiens ADP-ribosylation factor guanine nucleotide-exchange factor 2 (brefedidin A-inhibited) (ARFGEF2), mRNA  NM_005992  Homo sapiens T-box 1 (TBX1), transcript variant B, mRNA  NM_00383  Homo sapiens ranslocase of inner mitochondrial membrane 17 homolog B (yeast) (TIMM17B), mRNA  NM_000385  Homo sapiens aquaporin 1 (channel-forming integral protein, 28kD) (AQP1), mRNA  NM_000385  Homo sapiens Ras protein-specific guanine nucleotide-releasing factor 1 (RASGRF1), mRNA  NM_000963  Homo sapiens Ras protein-specific guanine nucleotide-releasing factor 1 (RASGRF1), mRNA  NM_000963  Homo sapiens grataglandin-endoperoxide synthase 2 (prostaglandin G/H synthase and cyclooxygenase) (PTGS2), mRNA  NM_002392  Homo sapiens proteasome (prosome, macropain) subunit, alpha type, 7 (PSMA7), mRNA  NM_003335  Homo sapiens low density lipoprotein receptor-related protein 5 (LRP5), mRNA  NM_004067  Homo sapiens dynein light chain 2 (Dle2), mRNA  NM_080677  Homo sapiens dynein light chain 2 (Dle2), mRNA  NM_080671  Homo sapiens synthal roll (KEH17739), mRNA  NM_080671  Homo sapiens potassium voltage-gated channel, Isk-related subfamily, gene 4 (KCNE4), mlRNA  NM_080667  Homo sapiens similar to RIKEN cDNA 2610030116 gene (MGC2541), mRNA  NM_080667  Homo sapiens similar to RIKEN cDNA 1110002C08 gene (MGC9564), mRNA  NM_080667  Homo sapiens similar to RIKEN cDNA 1110002C08 gene (MGC9564), mRNA  NM_080669  Homo sapiens similar to RIKEN cDNA 1810022F11 gene (MGC14289), mRNA  NM_080661  Homo sapiens similar to RIKEN cDNA 1810022F11 gene (MGC14289), mRNA  NM_080665  Homo sapiens similar to RIKEN cDNA 1800014N16 gene (MGC14289), mRNA  NM_080665  Homo sapiens similar to RIKEN cDNA 1800014N16 gene (MGC14289), mRNA  NM_080667  Homo sapiens similar to RIKEN cDNA 1800014N16 gene (MGC15407), mRNA  NM_080669  Homo sapiens similar to RIKEN cDNA 1800014N16 gene (MGC14289), mRNA  NM_080661  Homo sapiens similar to RIKEN cDNA 373052B13 gene (MGC15337), mRNA  NM_080665  Homo sapiens similar to RIKEN cDNA 373052B	NM 006335	Homo canions travels
NM_006420 Homo sapiens ADP-ribosylation factor guanine nucleotide-exchange factor 2 (brefeldin A-inhibited) (ARFGEF2), mRNA  NM_005834 Homo sapiens T-box 1 (TBX1), transcript variant B, mRNA  Homo sapiens translocase of inner mitochondrial membrane 17 homolog B (yeast) (TIMM17B), mRNA  NM_000385 Homo sapiens as protein-specific guanine nucleotide-releasing factor 1 (RASGRF1), mRNA  NM_0002891 Homo sapiens Ras protein-specific guanine nucleotide-releasing factor 1 (RASGRF1), mRNA  NM_000963 Homo sapiens prostaglandin-endoperoxide synthase 2 (prostaglandin G/H synthases and cyclooxygenase) (PTGS2), mRNA  NM_002792 Homo sapiens proteasome (prosome, macropain) subunit, alpha type, 7 (PSMA7), mRNA  NM_002335 Homo sapiens proteasome (prosome, macropain) subunit, alpha type, 7 (PSMA7), mRNA  NM_001402 Homo sapiens eukaryotic translation elongation factor 1 alpha 1 (EEF1A1), mRNA  NM_080671 Homo sapiens dynein light chain 2 (Dlc2), mRNA  NM_080672 Homo sapiens dynein light chain 2 (Dlc2), mRNA  NM_080671 Homo sapiens similar to RIKEN cDNA 2610030J16 gene (MGC2541), mRNA  NM_080669 Homo sapiens similar to RIKEN cDNA 4931428D14 gene (MGC15407), mRNA  NM_080669 Homo sapiens similar to RIKEN cDNA 1110002C08 gene (MGC9564), mRNA  NM_080660 Homo sapiens similar to RIKEN cDNA 181002EF11 gene (MGC19604), mRNA  NM_080664 Homo sapiens similar to RIKEN cDNA 181002EF11 gene (MGC19604), mRNA  NM_080665 Homo sapiens similar to RIKEN cDNA 181002GF1 gene (MGC19604), mRNA  NM_080666 Homo sapiens similar to RIKEN cDNA 18003G06 gene (MGC14289), mRNA  NM_080665 Homo sapiens similar to RIKEN cDNA 18003G06 gene (MGC14839), mRNA  NM_080665 Homo sapiens similar to RIKEN cDNA 18003G06 gene (MGC14839), mRNA  NM_080666 Homo sapiens similar to RIKEN cDNA 18003G06 gene (MGC14839), mRNA  NM_080667 Homo sapiens similar to RIKEN cDNA 18003G06 gene (MGC14839), mRNA  NM_080669 Homo sapiens similar to RIKEN cDNA 18003G06 gene (MGC14839), mRNA  NM_080660 Homo sapiens similar to RIKEN cDNA 18003G06 gene (MGC14839), mRNA  NM_080661 Homo sapiens similar to		and the state of t
(brefeldim A-inhibited) (ARFGBF2), mRNA  NM 005992  Homo sapiens T-box 1 (TBX1), transcript variant B, mRNA  Homo sapiens T-box 1 (TBX1), transcript variant B, mRNA  NM 000385  Homo sapiens ranislocase of inner mitochondrial membrane 17 homolog B  (yeast) (TIMM17B), mRNA  NM 000385  Homo sapiens aquaporin 1 (channel-forming integral protein, 28kD) (AQP1), mRNA  NM 002891  Homo sapiens Ras protein-specific guanine nucleotide-releasing factor 1  (RASGRF1), mRNA  NM 000963  Homo sapiens proteasglandin-endoperoxide synthase 2 (prostaglandin G/H synthase and cyclooxygenase) (PTGS2), mRNA  Homo sapiens proteasome (prosome, macropain) subunit, alpha type, 7  (PSMA7), mRNA  NM 002335  Homo sapiens low density lipoprotein receptor-related protein 5 (LRP5), mRNA  NM 001402  Homo sapiens density lipoprotein receptor-related protein 5 (LRP5), mRNA  NM 080677  Homo sapiens dynein light chain 2 (Dlc2), mRNA  NM 080671  Homo sapiens dynein light chain 2 (Dlc2), mRNA  NM 080671  Homo sapiens godssium voltage-gated channel, Isk-related subfamily, gene 4  (KCNE4), mRNA  NM 080667  Homo sapiens similar to RIKEN cDNA 2610030116 gene (MGC2541), mRNA  NM 080669  Homo sapiens similar to RIKEN cDNA 2810030116 gene (MGC9564), mRNA  NM 080660  Homo sapiens similar to RIKEN cDNA 4931428D14 gene (MGC15407), mRNA  NM 080660  Homo sapiens similar to RIKEN cDNA 1810022F11 gene (MGC19604), mRNA  NM 080661  Homo sapiens similar to RIKEN cDNA 1810022F11 gene (MGC19604), mRNA  NM 080655  Homo sapiens similar to RIKEN cDNA 1810022F11 gene (MGC19604), mRNA  NM 080654  Homo sapiens similar to RIKEN cDNA 1810022F11 gene (MGC19604), mRNA  NM 080655  Homo sapiens similar to RIKEN cDNA 1810022F11 gene (MGC19604), mRNA  NM 080656  Homo sapiens similar to RIKEN cDNA 1810022F11 gene (MGC19604), mRNA  Homo sapiens similar to RIKEN cDNA 1810022F11 gene (MGC19604), mRNA  Homo sapiens similar to RIKEN cDNA 1810022F11 gene (MGC19604), mRNA  Homo sapiens similar to RIKEN cDNA 5730528L13 gene (MGC15397), mRNA  Homo sapiens similar to RIKEN cDNA 5730528L13 gene (	NM 006420	
NM   005992   Homo sapiens T-box I (TBXI), transcript variant B, mRNA   Homo sapiens translocase of inner mitochondrial membrane 17 homolog B (yeast) (TIMM17B), mRNA   Homo sapiens aquaporin I (channel-forming integral protein, 28kD) (AQPI), mRNA   Homo sapiens Ras protein-specific guanine nucleotide-releasing factor I (RASGRFI), mRNA   Homo sapiens prostaglandin-endoperoxide synthase 2 (prostaglandin G/H synthases and cyclooxygenase) (PTGS2), mRNA   Homo sapiens proteasome (prosome, macropain) subunit, alpha type, 7 (PSMA7), mRNA   Homo sapiens proteasome (prosome, macropain) subunit, alpha type, 7 (PSMA7), mRNA   Homo sapiens broteasome (prosome, macropain) subunit, alpha type, 7 (PSMA7), mRNA   Homo sapiens own density lipoprotein receptor-related protein 5 (LRP5), mRNA   MN   001402   Homo sapiens own density lipoprotein receptor-related protein 5 (LRP5), mRNA   MN   080677   Homo sapiens dynein light chain 2 (Dlc2), mRNA   Homo sapiens oynein light chain 2 (Dlc2), mRNA   MN   080671   Homo sapiens oynein light chain 2 (Dlc2), mRNA   MN   080671   Homo sapiens potassium voltage-gated channel, lsk-related subfamily, gene 4 (KCNE4), mRNA   MN   080667   Homo sapiens similar to RIKEN cDNA 2610030116 gene (MGC2541), mRNA   MN   080667   Homo sapiens similar to RIKEN cDNA 1110002C08 gene (MGC9564), mRNA   MN   080667   Homo sapiens similar to RIKEN cDNA 181002F11 gene (MGC1407), mRNA   MN   080664   Homo sapiens similar to RIKEN cDNA 181002F11 gene (MGC19604), mRNA   MN   080665   Homo sapiens similar to RIKEN cDNA 181002F11 gene (MGC14289), mRNA   MN   080665   Homo sapiens similar to RIKEN cDNA 181002F11 gene (MGC14289), mRNA   MN   080665   Homo sapiens similar to RIKEN cDNA 181002F11 gene (MGC14289), mRNA   MN   080665   Homo sapiens similar to RIKEN cDNA 181002F11 gene (MGC14289), mRNA   MN   080665   Homo sapiens similar to RIKEN cDNA 181002F11 gene (MGC14289), mRNA   MN   080655   Homo sapiens similar to RIKEN cDNA 181002F11 gene (MGC19377), mRNA   MN   080651   Homo sapiens similar to RIKEN cDNA 181002	-1111_000420	
Mono Sapiens 1-box 1 (TBX1), transcript variant B, mRNA   Mono Sapiens translocase of inner mitochondrial membrane 17 homolog B   Mono sapiens similar to RIKEN cDNA 193118G17 gene (MGC1541), mRNA   Mono Sapiens similar to RIKEN cDNA 19305065   Homo sapiens similar to RIKEN cDNA 19305078   Homo Sapiens similar to RIKEN cDNA 19305078   Homo Sapiens similar to RIKEN cDNA 193050791, mRNA   Mono Sapiens similar to RIKEN cDNA 19305078   Homo Sapiens simi	NM 005000	
NM_000385 NM_000385 NM_000385 NM_000385 NM_000385 NM_000385 NM_0003891 NM_002891 Homo sapiens Ras protein-specific guanine nucleotide-releasing factor 1 (RASGRFI), mRNA NM_000963 Homo sapiens Prostaglandin-endoperoxide synthase 2 (prostaglandin G/H synthase and cyclooxygenase) (PTGS2), mRNA NM_000963 NM_002792 NM_002792 Homo sapiens proteasome (prosome, macropain) subunit, alpha type, 7 (PSMA7), mRNA NM_002335 NM_002335 Homo sapiens low density lipoprotein receptor-related protein 5 (LRP5), mRNA NM_001402 Homo sapiens eukaryotic translation elongation factor 1 alpha 1 (EEF1A1), mRNA NM_080677 Homo sapiens dynein light chain 2 (Dlc2), mRNA NM_080677 Homo sapiens dynein light chain 2 (Dlc2), mRNA NM_080671 Homo sapiens similar to RIKEN cDNA 2610030J16 gene (MGC2541), mRNA NM_080670 Homo sapiens similar to RIKEN cDNA 1110002C08 gene (MGC2541), mRNA NM_080660 Homo sapiens similar to RIKEN cDNA 4931428D14 gene (MGC194407), mRNA NM_080661 Homo sapiens similar to RIKEN cDNA 4930578F06 gene (MGC19604), mRNA NM_080662 Homo sapiens similar to RIKEN cDNA 1810022F11 gene (MGC19604), mRNA NM_080664 Homo sapiens similar to RIKEN cDNA 2310030G06 gene (MGC281), mRNA NM_080665 Homo sapiens similar to RIKEN cDNA 1810022F11 gene (MGC19604), mRNA NM_080665 Homo sapiens similar to RIKEN cDNA 1810022F11 gene (MGC19604), mRNA NM_080665 Homo sapiens similar to RIKEN cDNA 1810022F11 gene (MGC19489), mRNA NM_080655 Homo sapiens similar to RIKEN cDNA 1810022F11 gene (MGC14289), mRNA NM_080655 Homo sapiens similar to RIKEN cDNA 1810027F11 gene (MGC14389), mRNA NM_080655 Homo sapiens similar to RIKEN cDNA 1810027F11 gene (MGC19341), mRNA NM_080655 Homo sapiens similar to RIKEN cDNA 1810027F11 gene (MGC19341), mRNA NM_080655 Homo sapiens similar to RIKEN cDNA 1810027F11 gene (MGC19341), mRNA NM_080655 Homo sapiens similar to RIKEN cDNA 1810027F11 gene (MGC19341), mRNA NM_080655 Homo sapiens similar to RIKEN cDNA 1810027F11 gene (MGC19341), mRNA NM_080655 Homo sapiens similar to RIKEN cDNA 1810027F11 gene (MGC19377), mRNA NM_080655 Homo sapi		riomo sapiens T-box 1 (TBX1) transcript vorient D. Dald
NM_000385 Homo sapiens aquaporin 1 (channel-forming integral protein, 28kD) (AQP1), mRNA NM_002891 Homo sapiens Ras protein-specific guanine nucleotide-releasing factor 1 (RASGRF1), mRNA NM_000963 Homo sapiens prostaglandin-endoperoxide synthase 2 (prostaglandin G/H synthase and cyclooxygenase) (PTGS2), mRNA NM_002792 Homo sapiens proteasome (prosome, macropain) subunit, alpha type, 7 (PSMA7), mRNA NM_002335 Homo sapiens low density lipoprotein receptor-related protein 5 (LRP5), mRN/2 NM_001402 Homo sapiens eukaryotic translation elongation factor 1 alpha 1 (EEF1A1), mRNA NM_080677 NM_080677 NM_080671 NM_080671 Homo sapiens dynein light chain 2 (Dlc2), mRNA NM_080671 Homo sapiens potassium voltage-gated channel, Isk-related subfamily, gene 4 (KCNE4), mRNA NM_080667 Homo sapiens similar to RIKEN cDNA 2610030J16 gene (MGC2541), mRNA NM_080667 Homo sapiens similar to RIKEN cDNA 2110002C08 gene (MGC9564), mRNA NM_080667 Homo sapiens similar to RIKEN cDNA 4931428D14 gene (MGC15407), mRN/2 NM_080665 Homo sapiens similar to RIKEN cDNA 4931628D14 gene (MGC19604), mRNA NM_080666 Homo sapiens similar to RIKEN cDNA 1810022F11 gene (MGC19604), mRNA NM_080669 Homo sapiens similar to RIKEN cDNA 1810022F11 gene (MGC19804), mRNA NM_080669 Homo sapiens similar to RIKEN cDNA 180022F11 gene (MGC14289), mRN/3 NM_080669 Homo sapiens similar to RIKEN cDNA 1800014N16 gene (MGC14289), mRN/4 NM_080659 Homo sapiens similar to RIKEN cDNA 5730528L13 gene (MGC19337), mRN/4 NM_080651 Homo sapiens similar to RIKEN cDNA 5730528L13 gene (MGC17337), mRN/4 NM_080653 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRN/4 NM_080654 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRN/4 NM_080654 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRN/4 NM_080654 Homo sapiens similar to RIKEN cDNA 5700578N08 gene (MGC15397), mRN/4 NM_080654 Homo sapiens similar to RIKEN cDNA 5700578N08 gene (MGC15397), mRN/4 NM_080654 Homo sapiens similar to RIKEN cDNA 5700578N08 gene (MGC15397), mRN/4 NM_080655 Homo sapiens s	1NIM_005834	fromo sapiens translocase of inner mitochondrial mombres, 171
Homo sapiens aquaporin 1 (channel-forming integral protein, 28kD) (AQP1), mRNA   Homo sapiens Ras protein-specific guanine nucleotide-releasing factor 1 (RASGRF1), mRNA   Homo sapiens proteasome (prosome, macropain) subunit, alpha type, 7 (PSMA7), mRNA   Homo sapiens proteasome (prosome, macropain) subunit, alpha type, 7 (PSMA7), mRNA   Homo sapiens low density lipoprotein receptor-related protein 5 (LRP5), mRNA   Homo sapiens cukaryotic translation elongation factor 1 alpha 1 (EEF1A1), mRNA   Homo sapiens dynein light chain 2 (Dlc2), mRNA   Homo sapiens eukaryotic translation elongation factor 1 alpha 1 (EEF1A1), mRNA   Homo sapiens dynein light chain 2 (Dlc2), mRNA   Homo sapiens dynein light chain 2 (Dlc2), mRNA   Homo sapiens operated channel, lsk-related subfamily, gene 4 (KCNE4), mRNA   Homo sapiens potassium voltage-gated channel, lsk-related subfamily, gene 4 (KCNE4), mRNA   Homo sapiens similar to RIKEN cDNA 2610030J16 gene (MGC2541), mRNA   MM 080667   Homo sapiens similar to RIKEN cDNA 2911428D14 gene (MGC15407), mRNA   MM 080665   Homo sapiens similar to RIKEN cDNA 4931428D14 gene (MGC19604), mRNA   MM 080664   Homo sapiens similar to RIKEN cDNA 49310578F06 gene (MGC19604), mRNA   MM 080665   Homo sapiens similar to RIKEN cDNA 1810022F11 gene (MGC19604), mRNA   MM 080660   Homo sapiens similar to RIKEN cDNA 2930500C14 gene (MGC14839), mRNA   MM 080651   Homo sapiens similar to RIKEN cDNA 3730528L13 gene (MGC14839), mRNA   MM 080653   Homo sapiens similar to RIKEN cDNA 3730578N08 gene (MGC15397), mRNA   Homo sapiens similar to RIKEN cDNA 3730578N08 gene (MGC15397), mRNA   Homo sapiens similar to RIKEN cDNA 3730578N08 gene (MGC15397), mRNA   Homo sapiens similar to RIKEN cDNA 3730578N08 gene (MGC15397), mRNA   Homo sapiens similar to RIKEN cDNA 3730578N08 gene (MGC15397), mRNA   Homo sapiens similar to RIKEN cDNA 3730578N08 gene (MGC15397), mRNA   Homo sapiens similar to RIKEN cDNA 3730578N08 gene (MGC15397), mRNA   Homo sapiens chromosome 20 open reading frame 160 (C20orf160), mRNA   Homo sapiens c	-	(J + wor) ( + 11 + 11 + 1 + 1 + 1 + 1 + 1 + 1 + 1
NM_002891  Homo sapiens Ras protein-specific guanine nucleotide-releasing factor 1  (RASGRF1), mRNA  NM_000963  Homo sapiens prostaglandin-endoperoxide synthase 2 (prostaglandin G/H synthase and cyclooxygenase) (PTGS2), mRNA  Homo sapiens proteasome (prosome, macropain) subunit, alpha type, 7  (PSMA7), mRNA  MM_002335  Homo sapiens low density lipoprotein receptor-related protein 5 (LRP5), mRNA  NM_001402  Homo sapiens eukaryotic translation elongation factor 1 alpha 1 (EEF1A1), mRNA  NM_080677  Homo sapiens dynein light chain 2 (Dic2), mRNA  NM_080672  Homo sapiens gyl4474 like (H17739), mRNA  NM_080671  Homo sapiens gyl4474 like (H17739), mRNA  NM_080670  Homo sapiens similar to RIKEN cDNA 2610030J16 gene (MGC2541), mRNA  NM_080667  Homo sapiens similar to RIKEN cDNA 1110002C08 gene (MGC2541), mRNA  NM_080669  Homo sapiens similar to RIKEN cDNA 4931428D14 gene (MGC15407), mRNA  NM_080661  Homo sapiens similar to RIKEN cDNA 4930578F06 gene (MGC9564), mRNA  NM_080662  Homo sapiens similar to RIKEN cDNA 1810022F11 gene (MGC19604), mRNA  NM_080663  Homo sapiens similar to RIKEN cDNA 181002F11 gene (MGC4281), mRNA  NM_080664  Homo sapiens similar to RIKEN cDNA 181002F11 gene (MGC14839), mRNA  NM_080665  Homo sapiens similar to RIKEN cDNA 2310030G06 gene (MGC14839), mRNA  NM_080666  Homo sapiens similar to RIKEN cDNA 2310030G06 gene (MGC14839), mRNA  NM_080657  Homo sapiens similar to RIKEN cDNA 4930578F08 gene (MGC14839), mRNA  NM_080658  Homo sapiens similar to RIKEN cDNA 2310030G06 gene (MGC14839), mRNA  NM_080659  Homo sapiens similar to RIKEN cDNA 5730528L13 gene (MGC17337), mRNA  NM_080651  Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC13397), mRNA  NM_080653  Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC13397), mRNA  NM_080654  Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC13397), mRNA  NM_080654  Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC13397), mRNA  NM_080665  Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC13397), mRNA  NM_080666  Homo sapiens similar to RI	NM_000385	Homo sapiens aquaporin 1 (channel-forming integral protein 2017) (A OP1)
(RASGRF1), mRNA  NM_000963  Homo sapiens prostaglandin-endoperoxide synthase 2 (prostaglandin G/H synthase and cyclooxygenase) (PTGS2), mRNA  NM_002792  Homo sapiens proteasome (prosome, macropain) subunit, alpha type, 7 (PSMA7), mRNA  NM_001402  Homo sapiens eukaryotic translation elongation factor 1 alpha 1 (EEF1A1), mRNA  NM_080677  Homo sapiens dynein light chain 2 (Dlc2), mRNA  NM_080671  Homo sapiens dynein light chain 2 (Dlc2), mRNA  NM_080671  Homo sapiens potassium voltage-gated channel, Isk-related subfamily, gene 4 (KCNE4), mRNA  NM_080670  Homo sapiens similar to RIKEN cDNA 2610030J16 gene (MGC2541), mRNA  NM_080667  Homo sapiens similar to RIKEN cDNA 1110002C08 gene (MGC9564), mRNA  NM_080667  Homo sapiens similar to RIKEN cDNA 8230118G17 gene (MGC15407), mRNA  NM_080667  Homo sapiens similar to RIKEN cDNA 4930578F06 gene (MGC19604), mRNA  NM_080664  Homo sapiens similar to RIKEN cDNA 1100014N16 gene (MGC14289), mRNA  NM_080665  Homo sapiens similar to RIKEN cDNA 1200014N16 gene (MGC14289), mRNA  NM_080669  Homo sapiens similar to RIKEN cDNA 1200014N16 gene (MGC14289), mRNA  NM_080669  Homo sapiens similar to RIKEN cDNA 1200014N16 gene (MGC14289), mRNA  NM_080669  Homo sapiens similar to RIKEN cDNA 1200014N16 gene (MGC14389), mRNA  NM_080669  Homo sapiens similar to RIKEN cDNA 130030G06 gene (MGC14389), mRNA  NM_080659  Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC13377), mRNA  NM_080651  Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA  NM_080652  Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA  NM_080661  Homo sapiens similar to RIKEN cDNA 1200014N16, mRNA  Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA  NM_080654  Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA  NM_080655  Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA  NM_080661  Homo sapiens similar to RIKEN cDNA 1200014N16, mRNA  Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA  NM_08061  Homo sapiens similar to RI		
NM_000963   Homo sapiens prostaglandin-endoperoxide synthase 2 (prostaglandin G/H synthase and cyclooxygenase) (PTGS2), mRNA	NM_002891	Homo sapiens Ras protein-specific quanine puologida and
NM_000963 Homo sapiens prostaglandin-endoperoxide synthase 2 (prostaglandin G/H synthase and cyclooxygenase) (PTGS2), mRNA  NM_002395 Homo sapiens proteasome (prosome, macropain) subunit, alpha type, 7  (PSMA7), mRNA  NM_002335 Homo sapiens low density lipoprotein receptor-related protein 5 (LRP5), mRNA  NM_001402 Homo sapiens eukaryotic translation elongation factor 1 alpha 1 (EEF1A1), mRNA  NM_080677 Homo sapiens dynein light chain 2 (Dic2), mRNA  NM_080678 Homo sapiens gy9H4T4 like (H17739), mRNA  NM_080679 Homo sapiens similar to RIKEN cDNA 2610030J16 gene (MGC2541), mRNA  NM_080660 Homo sapiens similar to RIKEN cDNA 2610030J16 gene (MGC2541), mRNA  NM_080661 Homo sapiens similar to RIKEN cDNA 4931428D14 gene (MGC15407), mRNA  NM_080662 Homo sapiens similar to RIKEN cDNA 4930578F06 gene (MGC19604), mRNA  NM_080664 Homo sapiens similar to RIKEN cDNA 1110002Z011 gene (MGC19604), mRNA  NM_080665 Homo sapiens similar to RIKEN cDNA 4930578F06 gene (MGC9912), mRNA  NM_080660 Homo sapiens similar to RIKEN cDNA 12100014N16 gene (MGC14289), mRNA  NM_080659 Homo sapiens similar to RIKEN cDNA 1200014N16 gene (MGC14289), mRNA  NM_080650 Homo sapiens similar to RIKEN cDNA 1200014N16 gene (MGC14289), mRNA  NM_080651 Homo sapiens similar to RIKEN cDNA 1200014N16 gene (MGC1439), mRNA  NM_080653 Homo sapiens similar to RIKEN cDNA 1200014N16 gene (MGC1439), mRNA  NM_080654 Homo sapiens similar to RIKEN cDNA 1200014N16 gene (MGC1439), mRNA  NM_080655 Homo sapiens similar to RIKEN cDNA 1200014N16 gene (MGC1337), mRNA  NM_080654 Homo sapiens similar to RIKEN cDNA 1200014N16 gene (MGC1337), mRNA  NM_080655 Homo sapiens similar to RIKEN cDNA 1200014N16 gene (MGC1337), mRNA  NM_080654 Homo sapiens similar to RIKEN cDNA 1200014N16 gene (MGC1337), mRNA  NM_080655 Homo sapiens similar to RIKEN cDNA 1200014N16 gene (MGC1337), mRNA  NM_080664 Homo sapiens similar to RIKEN cDNA 1200014N16 gene (MGC1337), mRNA  NM_080655 Homo sapiens similar to RIKEN cDNA 1200014N16 gene (MGC1337), mRNA  NM_080666 Homo sapiens similar to RIKEN cDNA 12000		
NM_002792	NM_000963	Homo sapiens prostaglandin-endonerovido gratha 2 (
rhomo sapiens proteasome (prosome, macropain) subunit, alpha type, 7 (PSMA7), mRNA  NM_001402 Homo sapiens low density lipoprotein receptor-related protein 5 (LRP5), mRNA mRNA NM_080671 Homo sapiens dynein light chain 2 (Dlc2), mRNA NM_080672 Homo sapiens dynein light chain 2 (Dlc2), mRNA NM_080671 Homo sapiens potassium voltage-gated channel, Isk-related subfamily, gene 4 (KCNE4), mRNA NM_080667 Homo sapiens similar to RIKEN cDNA 2610030J16 gene (MGC2541), mRNA NM_080669 Homo sapiens similar to RIKEN cDNA 1110002C08 gene (MGC9564), mRNA NM_080667 Homo sapiens similar to RIKEN cDNA 4931428D14 gene (MGC15407), mRNA NM_080667 Homo sapiens similar to RIKEN cDNA B230118G17 gene (MGC19604), mRNA NM_080664 Homo sapiens similar to RIKEN cDNA 4930578F06 gene (MGC9912), mRNA NM_080665 Homo sapiens similar to RIKEN cDNA 1810022F11 gene (MGC4281), mRNA NM_080669 Homo sapiens similar to RIKEN cDNA 2310030G06 gene (MGC14289), mRNA NM_080659 Homo sapiens similar to RIKEN cDNA 2310030G06 gene (MGC14289), mRNA NM_080659 Homo sapiens similar to RIKEN cDNA 5730528L13 gene (MGC17337), mRNA NM_080651 Homo sapiens similar to RIKEN cDNA 5730528L13 gene (MGC17337), mRNA NM_080653 Homo sapiens similar to RIKEN cDNA 5730528L13 gene (MGC17337), mRNA NM_080654 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA NM_080655 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA NM_080654 Homo sapiens regulator of G-protein signalling 6 (RGS6), mRNA NM_042775 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA NM_080660 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA NM_080661 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA NM_080661 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA NM_080661 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA NM_080601 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA NM_080601 Homo sapiens sentomosome 20 open reading frame 112 (C20orf160), mRNA NM_080601 Homo sapiens chro		synthase and cyclogygenase) (PTGS2) while
(PSMA7), miRNA  NM_002335  Homo sapiens low density lipoprotein receptor-related protein 5 (LRP5), miRNA  miRNA  NM_001402  Homo sapiens low density lipoprotein receptor-related protein 5 (LRP5), miRNA  miRNA  NM_080677  Homo sapiens dynein light chain 2 (Dlc2), miRNA  NM_080672  Homo sapiens potassium voltage-gated channel, Isk-related subfamily, gene 4 (KCNE4), miRNA  NM_080671  Homo sapiens similar to RIKEN cDNA 2610030J16 gene (MGC2541), miRNA  NM_080669  Homo sapiens similar to RIKEN cDNA 1110002C08 gene (MGC9564), miRNA  NM_080669  Homo sapiens similar to RIKEN cDNA 4931428D14 gene (MGC15407), miRNA  NM_080665  Homo sapiens similar to RIKEN cDNA 4930578F06 gene (MGC9912), miRNA  NM_080660  Homo sapiens similar to RIKEN cDNA 1200014N16 gene (MGC4281), miRNA  NM_080659  Homo sapiens similar to RIKEN cDNA 1200014N16 gene (MGC14289), miRNA  NM_080659  Homo sapiens similar to RIKEN cDNA 2310030G06 gene (MGC1489), miRNA  NM_080659  Homo sapiens similar to RIKEN cDNA 2310030G06 gene (MGC1489), miRNA  NM_080659  Homo sapiens similar to RIKEN cDNA 5730528L13 gene (MGC1489), miRNA  NM_080650  Homo sapiens similar to RIKEN cDNA 493050C14 gene (MGC1337), miRNA  NM_080651  Homo sapiens similar to RIKEN cDNA 493050C14 gene (MGC1337), miRNA  NM_080652  Homo sapiens similar to RIKEN cDNA 5730528L13 gene (MGC1337), miRNA  NM_080654  Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC1337), miRNA  NM_080655  Homo sapiens similar to RIKEN cDNA 493050C14 gene (MGC9341), miRNA  NM_080650  Homo sapiens similar to RIKEN cDNA 493050C14 gene (MGC9341), miRNA  NM_080651  Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC1337), miRNA  NM_080652  Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC1337), miRNA  NM_080654  Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC1337), miRNA  NM_080655  Homo sapiens chromosome 20 open reading frame 160 (C20orf160), miRNA  NM_080661  Homo sapiens chromosome 20 open reading frame 160 (C20orf160), miRNA  NM_080661  Homo sapiens chromosome 20 open reading frame 162 (C20orf162),	NM_002792	Homo saniens protessome (prosesses)
NM	_	(PSMA7) mRNA
NM_001402	NM 002335	
mRNA  NM 080677  Homo sapiens dynein light chain 2 (Dlc2), mRNA  NM 080672  Homo sapiens Q9H4T4 like (H17739), mRNA  NM 080671  Homo sapiens potassium voltage-gated channel, Isk-related subfamily, gene 4 (KCNE4), mRNA  NM 080670  Homo sapiens similar to RIKEN cDNA 2610030J16 gene (MGC2541), mRNA  NM 080669  Homo sapiens similar to RIKEN cDNA 1110002C08 gene (MGC9564), mRNA  NM 080667  Homo sapiens similar to RIKEN cDNA 4931428D14 gene (MGC15407), mRNA  NM 080668  Homo sapiens similar to RIKEN cDNA 4930578F06 gene (MGC9912), mRNA  NM 080664  Homo sapiens similar to RIKEN cDNA 1810022F11 gene (MGC4281), mRNA  NM 080659  Homo sapiens similar to RIKEN cDNA 1200014N16 gene (MGC14289), mRNA  NM 080650  Homo sapiens similar to RIKEN cDNA 2310030G06 gene (MGC14839), mRNA  NM 080651  Homo sapiens similar to RIKEN cDNA 5730528L13 gene (MGC17337), mRNA  NM 080653  Homo sapiens similar to RIKEN cDNA 4930578N08 gene (MGC17337), mRNA  NM 080654  Homo sapiens similar to RIKEN cDNA 4930500C14 gene (MGC9341), mRNA  NM 080655  Homo sapiens similar to RIKEN cDNA 4930500C14 gene (MGC9341), mRNA  NM 080656  Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA  NM 080650  Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA  NM 080655  Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA  NM 080656  Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA  NM 080656  Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA  NM 080655  Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA  NM 080656  Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA  NM 080650  Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA  NM 080651  Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA  NM 080651  Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA  NM 080601  Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC17337), mRNA  NM 080601  Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC17337),		Homo sepiens low density lipoprotein receptor-related protein 5 (LRP5), mRNA
NM 080677 NM 080672 NM 080672 NM 080672 NM 080672 NM 080673 NM 080671 NM 080671 NM 080671 NM 080670 NM 080660 NM 080669 NM 080669 NM 080667 NM 080660 NM 080661 NM 080660 NM 080660 NM 080661 NM 080662 NM 080660 NM 080662 NM 080660 NM 080650 NM 080650 NM 080650 NM 080650 NM 080650 NM 080651 NM 080651 NM 080652 NM 080652 NM 080653 NM 080653 NM 080654 NM 080653 NM 080654 NM 080655 NM 080655 NM 080655 NM 080655 NM 080656 NM 080656 NM 080656 NM 080656 NM 080657 NM 080657 NM 080658 NM 080659 NM 080659 NM 080650 NM 080650 NM 080650 NM 080651 NM 080651 NM 080652 NM 080652 NM 080653 NM 080654 NM 080655 NM 080655 NM 080655 NM 080656 NM 080656 NM 080656 NM 080656 NM 080657 NM 080657 NM 080658 NM 080659 NM 080659 NM 080659 NM 080659 NM 080650 NM 080650 NM 080650 NM 080651 NM 080651 NM 080652 NM 080652 NM 080653 NM 080654 NM 080655 NM 080655 NM 080655 NM 080656 NM 080666 NM 080		and suproiss cural your utilistation elongation factor 1 alpha 1 (CCC) 41)
NM 080672 Homo sapiens Q9H4T4 like (H17739), mRNA  NM 080671 Homo sapiens potassium voltage-gated channel, Isk-related subfamily, gene 4 (KCNE4), mRNA  NM 080669 Homo sapiens similar to RIKEN cDNA 2610030J16 gene (MGC2541), mRNA NM 080667 Homo sapiens similar to RIKEN cDNA 1110002C08 gene (MGC9564), mRNA NM 080667 Homo sapiens similar to RIKEN cDNA 4931428D14 gene (MGC15407), mRNA NM 080665 Homo sapiens similar to RIKEN cDNA 4930578F06 gene (MGC9904), mRNA NM 080660 Homo sapiens similar to RIKEN cDNA 1810022F11 gene (MGC19604), mRNA NM 080660 Homo sapiens similar to RIKEN cDNA 1810022F11 gene (MGC4281), mRNA NM 080650 Homo sapiens similar to RIKEN cDNA 1200014N16 gene (MGC14289), mRNA NM 080651 Homo sapiens similar to RIKEN cDNA 2310030G06 gene (MGC14839), mRNA NM 080655 Homo sapiens similar to RIKEN cDNA 5730528L13 gene (MGC17337), mRNA NM 080654 Homo sapiens similar to RIKEN cDNA 5730528L13 gene (MGC17337), mRNA NM 080655 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC19341), mRNA NM 080654 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC19341), mRNA NM 080655 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC19341), mRNA NM 080656 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC19341), mRNA NM 080656 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA NM 080656 Homo sapiens segulator of G-protein signalling 6 (RGS6), mRNA Homo sapiens BRIS binding protein (BRI3BP), mRNA Homo sapiens BRIS binding protein (BRI3BP), mRNA Homo sapiens chromosome 20 open reading frame 160 (C20orf160), mRNA NM 08061 Homo sapiens chromosome 20 open reading frame 112 (C20orf112), mRNA NM 080603 Homo sapiens chromosome 20 open reading frame 102 (C20orf102), mRNA NM 080603 Homo sapiens chromosome 20 open reading frame 102 (C20orf102), mRNA NM 080604 Homo sapiens chromosome 20 open reading frame 102 (C20orf102), mRNA NM 080604 Homo sapiens chromosome 20 open reading frame 102 (C20orf102), mRNA NM 080604 Homo sapiens chromosome 20 open reading frame 103 (C20orf102), mRNA NM 080604 Homo sapiens	NM 080677	
NM_080671 Homo sapiens potassium voltage-gated channel, Isk-related subfamily, gene 4 (KCNE4), mRNA  NM_080670 Homo sapiens similar to RIKEN cDNA 2610030J16 gene (MGC2541), mRNA NM_080669 Homo sapiens similar to RIKEN cDNA 1110002C08 gene (MGC9564), mRNA NM_080667 Homo sapiens similar to RIKEN cDNA 4931428D14 gene (MGC15407), mRNA NM_080665 Homo sapiens similar to RIKEN cDNA B230118G17 gene (MGC19604), mRNA  NM_080664 Homo sapiens similar to RIKEN cDNA 4930578F06 gene (MGC9912), mRNA NM_080662 Homo sapiens similar to RIKEN cDNA 1810022F11 gene (MGC4281), mRNA NM_080669 Homo sapiens similar to RIKEN cDNA 1810022F11 gene (MGC4281), mRNA NM_080660 Homo sapiens similar to RIKEN cDNA 1200014N16 gene (MGC14289), mRNA NM_080657 Homo sapiens similar to RIKEN cDNA 2310030G06 gene (MGC14839), mRNA NM_080657 Homo sapiens similar to RIKEN cDNA 5730528L13 gene (MGC17337), mRNA NM_080654 Homo sapiens similar to RIKEN cDNA 4930500C14 gene (MGC9341), mRNA NM_080655 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA NM_080650 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA NM_080651 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA NM_080652 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA NM_080654 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA NM_080655 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA NM_080656 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA NM_080650 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA NM_080661 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC17337), mRNA NM_080665 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC14839), mRNA NM_080666 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC14839), mRNA NM_080666 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC14839), mRNA NM_080666 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC17337), mRNA NM_080666 Homo sapiens chromosome 20 open reading frame 102 (C20orf102), mRNA		Homo sapiens dynein light chain 2 (Dlc2), mRNA
Homo sapiens potassium voltage-gated channel, Isk-related subfamily, gene 4 (KCNE4), mRNA  NM_080660 Homo sapiens similar to RIKEN cDNA 2610030J16 gene (MGC2541), mRNA NM_080667 Homo sapiens similar to RIKEN cDNA 1110002C08 gene (MGC9564), mRNA NM_080667 Homo sapiens similar to RIKEN cDNA 4931428D14 gene (MGC15407), mRNA NM_080665 Homo sapiens similar to RIKEN cDNA 4930578F06 gene (MGC9912), mRNA NM_080664 Homo sapiens similar to RIKEN cDNA 4930578F06 gene (MGC9912), mRNA NM_080665 Homo sapiens similar to RIKEN cDNA 1810022F11 gene (MGC4281), mRNA NM_080660 Homo sapiens similar to RIKEN cDNA 1200014N16 gene (MGC14289), mRNA NM_080657 Homo sapiens similar to RIKEN cDNA 2310030G06 gene (MGC14289), mRNA NM_080657 Homo sapiens vipirin (cig5), mRNA NM_080654 Homo sapiens vipirin (cig5), mRNA NM_080655 Homo sapiens similar to RIKEN cDNA 5730528L13 gene (MGC14337), mRNA NM_080654 Homo sapiens RIKEN cDNA 4930500C14 gene (MGC9341), mRNA NM_080655 Homo sapiens similar to RIKEN cDNA 4930500C14 gene (MGC9341), mRNA NM_080654 Homo sapiens regulator of G-protein signalling 6 (RGS6), mRNA NM_04240 Homo sapiens semin 6 (GEMIN6), mRNA NM_042475 Homo sapiens gemin 6 (GEMIN6), mRNA NM_080626 Homo sapiens shrding protein (BRI3BP), mRNA NM_080616 Homo sapiens chromosome 20 open reading frame 160 (C20orf160), mRNA NM_080616 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA NM_080617 Homo sapiens chromosome 20 open reading frame 102 (C20orf162), mRNA NM_080618 Homo sapiens chromosome 20 open reading frame 102 (C20orf162), mRNA NM_080619 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA NM_08061 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA NM_08061 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA NM_08061 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA NM_08061 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA NM_08061 Homo sapiens chromosome 20 open reading frame 164 (C20orf162), mRNA NM_08061 Homo sapiens chromosome 20		Tiono sapiens O9H414 like (H17730) mDNA
NM_080667 Homo sapiens similar to RIKEN cDNA 2610030J16 gene (MGC2541), mRNA NM_080669 Homo sapiens similar to RIKEN cDNA 1110002C08 gene (MGC9564), mRNA NM_080665 Homo sapiens similar to RIKEN cDNA 4931428D14 gene (MGC15407), mRNA NM_080665 Homo sapiens similar to RIKEN cDNA B230118G17 gene (MGC19604), mRNA NM_080664 Homo sapiens similar to RIKEN cDNA 4930578F06 gene (MGC9912), mRNA NM_080662 Homo sapiens similar to RIKEN cDNA 1810022F11 gene (MGC4281), mRNA NM_080659 Homo sapiens similar to RIKEN cDNA 1200014N16 gene (MGC14289), mRNA NM_080657 Homo sapiens similar to RIKEN cDNA 2310030G06 gene (MGC14839), mRNA NM_080657 Homo sapiens similar to RIKEN cDNA 5730528L13 gene (MGC17337), mRNA NM_080654 Homo sapiens similar to RIKEN cDNA 5730528L13 gene (MGC17337), mRNA NM_080655 Homo sapiens similar to RIKEN cDNA 4930500C14 gene (MGC9341), mRNA NM_080654 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA NM_080655 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA NM_080650 Homo sapiens regulator of G-protein signalling 6 (RGS6), mRNA NM_0424775 Homo sapiens FabG (beta-ketoacyl-[acyl-carrier-protein] reductase, E coli) like (E. coli) (FABGL), mRNA NM_080626 Homo sapiens BRI3 binding protein (BRI3BP), mRNA NM_080616 Homo sapiens chromosome 20 open reading frame 160 (C20orf160), mRNA NM_080601 Homo sapiens chromosome 20 open reading frame 112 (C20orf112), mRNA NM_080603 Homo sapiens chromosome 20 open reading frame 102 (C20orf102), mRNA NM_080603 Homo sapiens chromosome 20 open reading frame 102 (C20orf102), mRNA NM_080603 Homo sapiens chromosome 20 open reading frame 102 (C20orf102), mRNA NM_080604 Homo sapiens chromosome 20 open reading frame 126 (C20orf102), mRNA NM_080604 Homo sapiens chromosome 20 open reading frame 126 (C20orf102), mRNA NM_080604 Homo sapiens chromosome 20 open reading frame 126 (C20orf102), mRNA NM_080604 Homo sapiens chromosome 20 open reading frame 126 (C20orf102), mRNA NM_080604 Homo sapiens cask-interacting protein-like 8 (OSBPL8), mRNA	1/1/1 _0000/1	Homo sapiens potassium voltage-gated channel Telegralet 1 1 5
NM 080669 Homo sapiens similar to RIKEN cDNA 1110002C08 gene (MGC2541), mRNA NM 080667 Homo sapiens similar to RIKEN cDNA 4931428D14 gene (MGC9564), mRNA Homo sapiens similar to RIKEN cDNA 4931428D14 gene (MGC15407), mRNA mRNA Homo sapiens similar to RIKEN cDNA 4930578F06 gene (MGC9912), mRNA NM 080662 Homo sapiens similar to RIKEN cDNA 4930578F06 gene (MGC9912), mRNA NM 080660 Homo sapiens similar to RIKEN cDNA 1810022F11 gene (MGC4281), mRNA NM 080659 Homo sapiens similar to RIKEN cDNA 1200014N16 gene (MGC14289), mRNA NM 080657 Homo sapiens similar to RIKEN cDNA 2310030G06 gene (MGC14839), mRNA NM 080657 Homo sapiens vipirin (cig5), mRNA Homo sapiens similar to RIKEN cDNA 5730528L13 gene (MGC17337), mRNA NM 080654 Homo sapiens similar to RIKEN cDNA 5730528L13 gene (MGC17337), mRNA NM 080654 Homo sapiens similar to RIKEN cDNA 4930500C14 gene (MGC9341), mRNA NM 080654 Homo sapiens similar to RIKEN cDNA 4930500C14 gene (MGC9341), mRNA Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA NM 080650 Homo sapiens regulator of G-protein signalling 6 (RGS6), mRNA Homo sapiens FabG (beta-ketoacyl-[acyl-carrier-protein] reductase, E coli) like (E. coli) (FABGL), mRNA Homo sapiens BRI3 binding protein (BRI3BP), mRNA Homo sapiens brinding protein (BRI3BP), mRNA Homo sapiens chromosome 20 open reading frame 160 (C20orf160), mRNA Homo sapiens chromosome 20 open reading frame 112 (C20orf112), mRNA Homo sapiens chromosome 20 open reading frame 102 (C20orf162), mRNA Homo sapiens chromosome 20 open reading frame 102 (C20orf162), mRNA Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA MM 020464 Ho	ND C OCCUPA	
NM_080667 Homo sapiens similar to RIKEN cDNA 4931428D14 gene (MGC15407), mRNA mRNA_080665 Homo sapiens similar to RIKEN cDNA 4930578F06 gene (MGC19604), mRNA_080662 Homo sapiens similar to RIKEN cDNA 4930578F06 gene (MGC19604), mRNA_080662 Homo sapiens similar to RIKEN cDNA 1810022F11 gene (MGC4281), mRNA_080660 Homo sapiens similar to RIKEN cDNA 1810022F11 gene (MGC4281), mRNA_080657 Homo sapiens similar to RIKEN cDNA 1200014N16 gene (MGC14289), mRNA_080657 Homo sapiens vipirin (cig5), mRNA_080655 Homo sapiens vipirin (cig5), mRNA_080655 Homo sapiens similar to RIKEN cDNA 5730528L13 gene (MGC17337), mRNA_080654 Homo sapiens NY-REN-41 antigen (NY-REN-41), mRNA_080655 Homo sapiens similar to RIKEN cDNA 4930500C14 gene (MGC9341), mRNA_080651 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA_080651 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA_080651 Homo sapiens regulator of G-protein signalling 6 (RGS6), mRNA_080654 Homo sapiens RISI binding protein (BRI3BP), mRNA_080626 Homo sapiens BRI3 binding protein (BRI3BP), mRNA_080616 Homo sapiens chromosome 20 open reading frame 102 (C20orf112), mRNA_080616 Homo sapiens chromosome 20 open reading frame 112 (C20orf112), mRNA_080616 Homo sapiens chromosome 20 open reading frame 102 (C20orf102), mRNA_0806067 Homo sapiens chromosome 20 open reading frame 102 (C20orf102), mRNA_0806060 Homo sapiens chromosome 20 open reading frame 102 (C20orf102), mRNA_0806060 Homo sapiens chromosome 20 open reading frame 102 (C20orf102), mRNA_1806060 Homo sapiens chromosome 20 open reading frame 105 (C20orf102), mRNA_1806060 Homo sapiens chromosome 20 open reading frame 105 (C20orf102), mRNA_1806060 Homo sapiens chromosome 20 open reading frame 106 (C20orf102), mRNA_1806060 Homo sapiens chromosome 20 open reading frame 106 (C20orf102), mRNA_1806060 Homo sapiens chromosome 20 open reading frame 106 (C20orf102), mRNA_1806060 Homo sapiens chromosome 20 open reading frame 106 (C20orf102), mRNA_1806060 Homo sapiens chromosome 20 open reading frame 106 (C20o		Homo sapiens similar to RIKEN cDNA 2610030J16 gene (MGC2541) PNA
NM_080665 Homo sapiens similar to RIKEN cDNA B230118G17 gene (MGC19604), mRNA  NM_080664 Homo sapiens similar to RIKEN cDNA 4930578F06 gene (MGC9912), mRNA NM_080662 Homo sapiens similar to RIKEN cDNA 1810022F11 gene (MGC4281), mRNA NM_080669 Homo sapiens similar to RIKEN cDNA 1200014N16 gene (MGC14289), mRNA NM_080659 Homo sapiens similar to RIKEN cDNA 2310030G06 gene (MGC14289), mRNA NM_080657 Homo sapiens vipirin (cig5), mRNA NM_080658 Homo sapiens similar to RIKEN cDNA 5730528L13 gene (MGC17337), mRNA NM_080654 Homo sapiens similar to RIKEN cDNA 5730528L13 gene (MGC17337), mRNA NM_080655 Homo sapiens similar to RIKEN cDNA 4930500C14 gene (MGC9341), mRNA NM_080654 Homo sapiens similar to RIKEN cDNA 4930500C14 gene (MGC9341), mRNA NM_080655 Homo sapiens regulator of G-protein signalling 6 (RG86), mRNA NM_04296 Homo sapiens FabG (beta-ketoacyl-[acyl-carrier-protein] reductase, E coli) like (E. coli) (FABGL), mRNA NM_080626 Homo sapiens gemin 6 (GEMIN6), mRNA NM_080627 Homo sapiens BRI3 binding protein (BRI3BP), mRNA NM_080616 Homo sapiens chromosome 20 open reading frame 160 (C20orf160), mRNA NM_080617 Homo sapiens chromosome 20 open reading frame 112 (C20orf112), mRNA NM_080607 Homo sapiens chromosome 20 open reading frame 102 (C20orf102), mRNA NM_080607 Homo sapiens chromosome 20 open reading frame 162 (C20orf102), mRNA NM_080607 Homo sapiens chromosome 20 open reading frame 162 (C20orf160), mRNA NM_080607 Homo sapiens chromosome 20 open reading frame 162 (C20orf160), mRNA NM_080607 Homo sapiens chromosome 20 open reading frame 162 (C20orf160), mRNA NM_080607 Homo sapiens chromosome 20 open reading frame 162 (C20orf160), mRNA NM_080607 Homo sapiens chromosome 20 open reading frame 162 (C20orf160), mRNA NM_080607 Homo sapiens chromosome 20 open reading frame 162 (C20orf160), mRNA NM_080607 Homo sapiens chromosome 20 open reading frame 162 (C20orf160), mRNA NM_080607 Homo sapiens chromosome 20 open reading frame 162 (C20orf160), mRNA NM_080608 Homo sapiens chromosome 20 open reading frame 160 (C20orf160), mRNA		
mRNA  NM 080664 Homo sapiens similar to RIKEN cDNA 4930578F06 gene (MGC9912), mRNA  NM 080662 Homo sapiens similar to RIKEN cDNA 1810022F11 gene (MGC9912), mRNA  NM 080669 Homo sapiens similar to RIKEN cDNA 1200014N16 gene (MGC14289), mRNA  NM 080659 Homo sapiens similar to RIKEN cDNA 2310030G06 gene (MGC14289), mRNA  NM 080657 Homo sapiens vipirin (cig5), mRNA  NM 080655 Homo sapiens similar to RIKEN cDNA 5730528L13 gene (MGC17337), mRNA  NM 080654 Homo sapiens similar to RIKEN cDNA 5730528L13 gene (MGC17337), mRNA  NM 080655 Homo sapiens similar to RIKEN cDNA 4930500C14 gene (MGC9341), mRNA  NM 080654 Homo sapiens similar to RIKEN cDNA 4930500C14 gene (MGC9341), mRNA  NM 080655 Homo sapiens regulator of G-protein signalling 6 (RGS6), mRNA  NM 04296 Homo sapiens FabG (beta-ketoacyl-[acyl-carrier-protein] reductase, E coli) like  (E. coli) (FABGL), mRNA  NM 080626 Homo sapiens gemin 6 (GEMIN6), mRNA  NM 080627 Homo sapiens chromosome 20 open reading frame 160 (C20orf160), mRNA  NM 080610 Homo sapiens chromosome 20 open reading frame 112 (C20orf112), mRNA  NM 080601 Homo sapiens chromosome 20 open reading frame 102 (C20orf102), mRNA  NM 080603 Homo sapiens chromosome 20 open reading frame 162 (C20orf102), mRNA  NM 080604 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA  NM 080605 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA  NM 080601 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA  NM 080603 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA  NM 080604 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA  NM 080605 Homo sapiens chromosome 20 open reading frame 163 (C20orf162), mRNA  NM 080604 Homo sapiens chromosome 20 open reading frame 164 (C20orf162), mRNA  NM 080605 Homo sapiens chromosome 20 open reading frame 164 (C20orf162), mRNA  NM 080606 Homo sapiens chromosome 20 open reading frame 164 (C20orf162), mRNA  NM 080606 Homo sapiens chromosome 20 open reading frame 164 (C20orf162), mRNA  NM 080606 Homo sapiens chr		Homo sapiens similar to RIKEN cDNA 4931428D14 gene (MCC15407) PNA
NM 080664 Homo sapiens similar to RIKEN cDNA 4930578F06 gene (MGC9912), mRNA NM 080662 Homo sapiens similar to RIKEN cDNA 1810022F11 gene (MGC4281), mRNA NM 080660 Homo sapiens similar to RIKEN cDNA 1200014N16 gene (MGC14289), mRNA NM 080659 Homo sapiens similar to RIKEN cDNA 2310030G06 gene (MGC14289), mRNA NM 080657 Homo sapiens vipirin (cig5), mRNA NM 080655 Homo sapiens similar to RIKEN cDNA 5730528L13 gene (MGC17337), mRNA NM 080654 Homo sapiens similar to RIKEN cDNA 5730528L13 gene (MGC17337), mRNA NM 080655 Homo sapiens similar to RIKEN cDNA 4930500C14 gene (MGC9341), mRNA NM 080654 Homo sapiens similar to RIKEN cDNA 4930500C14 gene (MGC9341), mRNA NM 080655 Homo sapiens regulator of G-protein signalling 6 (RG86), mRNA NM 04296 Homo sapiens FabG (beta-ketoacyl-[acyl-carrier-protein] reductase, E coli) like (E. coli) (FABGL), mRNA NM 080626 Homo sapiens BRI3 binding protein (BR13BP), mRNA NM 080626 Homo sapiens chromosome 20 open reading frame 160 (C20orf160), mRNA NM 080616 Homo sapiens DOS/Gab family member 3 (GAB3), mRNA NM 080607 Homo sapiens chromosome 20 open reading frame 102 (C20orf102), mRNA NM 080603 Homo sapiens chromosome 20 open reading frame 162 (C20orf102), mRNA NM 080603 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA NM 030815 Homo sapiens chromosome 20 open reading frame 126 (C20orf162), mRNA NM 030815 Homo sapiens chromosome 20 open reading frame 126 (C20orf126), mRNA NM 030815 Homo sapiens chromosome 20 open reading frame 126 (C20orf162), mRNA NM 030814 Homo sapiens chromosome 20 open reading frame 126 (C20orf162), mRNA NM 030815 Homo sapiens chromosome 20 open reading frame 126 (C20orf162), mRNA NM 030816 Homo sapiens chromosome 20 open reading frame 126 (C20orf162), mRNA NM 030817 Homo sapiens chromosome 20 open reading frame 126 (C20orf162), mRNA NM 030816 Homo sapiens chromosome 20 open reading frame 126 (C20orf162), mRNA NM 030817 Homo sapiens chromosome 20 open reading frame 126 (C20orf162), mRNA NM 030816 Homo sapiens chromosome 20 open reading frame 126 (C20	NM_080665	Homo sapiens similar to RIKEN cDNA R230118G17 care (MGC13407), mRNA
MM_080662 Homo sapiens similar to RIKEN cDNA 1810022F11 gene (MGC9912), mRNA NM_080660 Homo sapiens similar to RIKEN cDNA 1810002F11 gene (MGC4281), mRNA NM_080659 Homo sapiens similar to RIKEN cDNA 1200014N16 gene (MGC14289), mRNA NM_080657 Homo sapiens similar to RIKEN cDNA 2310030G06 gene (MGC14839), mRNA NM_080655 Homo sapiens similar to RIKEN cDNA 5730528L13 gene (MGC17337), mRNA NM_080654 Homo sapiens similar to RIKEN cDNA 5730528L13 gene (MGC17337), mRNA NM_080655 Homo sapiens similar to RIKEN cDNA 4930500C14 gene (MGC9341), mRNA NM_080652 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA NM_080652 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA NM_080654 Homo sapiens regulator of G-protein signalling 6 (RGS6), mRNA NM_080655 Homo sapiens FabG (beta-ketoacy1-[acy1-carrier-protein] reductase, E coli) like (E. coli) (FABGL), mRNA NM_080626 Homo sapiens gemin 6 (GEMIN6), mRNA NM_080626 Homo sapiens BRI3 binding protein (BRI3BP), mRNA NM_080626 Homo sapiens chromosome 20 open reading frame 160 (C20orf160), mRNA NM_080616 Homo sapiens chromosome 20 open reading frame 112 (C20orf112), mRNA NM_080617 Homo sapiens chromosome 20 open reading frame 102 (C20orf162), mRNA NM_080603 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA NM_080615 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA NM_080616 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA NM_080617 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA NM_080603 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA NM_080615 Homo sapiens chromosome 20 open reading frame 164 (C20orf162), mRNA NM_080616 Homo sapiens chromosome 20 open reading frame 165 (C20orf162), mRNA NM_080617 Homo sapiens chromosome 20 open reading frame 165 (C20orf162), mRNA NM_080618 Homo sapiens chromosome 20 open reading frame 166 (C20orf162), mRNA NM_080618 Homo sapiens chromosome 20 open reading frame 166 (C20orf162), mRNA NM_080618 Homo sapiens chromosome 20 open		
MM_080660 Homo sapiens similar to RIKEN cDNA 1200014N16 gene (MGC14289), mRNA NM_080659 Homo sapiens similar to RIKEN cDNA 2310030G06 gene (MGC14839), mRNA NM_080657 Homo sapiens vipirin (cig5), mRNA NM_080655 Homo sapiens similar to RIKEN cDNA 2310030G06 gene (MGC14839), mRNA NM_080654 Homo sapiens similar to RIKEN cDNA 5730528L13 gene (MGC17337), mRNA NM_080653 Homo sapiens NY-REN-41 antigen (NY-REN-41), mRNA NM_080652 Homo sapiens similar to RIKEN cDNA 4930500C14 gene (MGC9341), mRNA NM_080652 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA NM_014234 Homo sapiens regulator of G-protein signalling 6 (RGS6), mRNA NM_024775 Homo sapiens FabG (beta-ketoacyl-[acyl-carrier-protein] reductase, E coli) like (E. coli) (FABGL), mRNA NM_080626 Homo sapiens BRI3 binding protein (BRI3BP), mRNA NM_080627 Homo sapiens chromosome 20 open reading frame 160 (C20orf160), mRNA NM_080616 Homo sapiens chromosome 20 open reading frame 112 (C20orf112), mRNA NM_080607 Homo sapiens chromosome 20 open reading frame 102 (C20orf102), mRNA NM_080603 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA NM_030815 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA NM_030815 Homo sapiens chromosome 20 open reading frame 162 (C20orf166), mRNA NM_020841 Homo sapiens chromosome 20 open reading frame 126 (C20orf166), mRNA NM_020841 Homo sapiens chromosome 20 open reading frame 162 (C20orf166), mRNA NM_020841 Homo sapiens chromosome 20 open reading frame 163 (C20orf166), mRNA NM_020841 Homo sapiens chromosome 20 open reading frame 164 (C20orf166), mRNA NM_020841 Homo sapiens chromosome 20 open reading frame 126 (C20orf166), mRNA NM_020841 Homo sapiens chromosome 20 open reading frame 1860 (C20orf166), mRNA NM_020841 Homo sapiens chromosome 20 open reading frame 1860 (C20orf166), mRNA NM_020841 Homo sapiens cask-interacting protein-like 8 (OSBPL8), mRNA		Homo sapiens similar to RIKEN CDNA 4020579796
MM_080659 Homo sapiens similar to RIKEN cDNA 2310030G06 gene (MGC14289), mRNA NM_080657 Homo sapiens vipirin (cig5), mRNA NM_080655 Homo sapiens similar to RIKEN cDNA 5730528L13 gene (MGC14839), mRNA NM_080654 Homo sapiens similar to RIKEN cDNA 5730528L13 gene (MGC17337), mRNA NM_080654 Homo sapiens NY-REN-41 antigen (NY-REN-41), mRNA NM_080655 Homo sapiens similar to RIKEN cDNA 4930500C14 gene (MGC9341), mRNA NM_080652 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA NM_080654 Homo sapiens regulator of G-protein signalling 6 (RGS6), mRNA NM_080655 Homo sapiens FabG (beta-ketoacyl-[acyl-carrier-protein] reductase, E coli) like (E. coli) (FABGL), mRNA NM_080626 Homo sapiens gemin 6 (GEMIN6), mRNA NM_080626 Homo sapiens chromosome 20 open reading frame 160 (C20orf160), mRNA NM_080616 Homo sapiens chromosome 20 open reading frame 112 (C20orf112), mRNA NM_080617 Homo sapiens chromosome 20 open reading frame 102 (C20orf102), mRNA NM_080603 Homo sapiens chromosome 20 open reading frame 102 (C20orf102), mRNA NM_080603 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA NM_080615 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA NM_080603 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA NM_080604 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA NM_080605 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA NM_080604 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA NM_080605 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA NM_080606 Homo sapiens chromosome 20 open reading frame 164 (C20orf164), mRNA NM_080606 Homo sapiens chromosome 20 open reading frame 165 (C20orf165), mRNA NM_080606 Homo sapiens chromosome 20 open reading frame 165 (C20orf166), mRNA NM_080606 Homo sapiens chromosome 20 open reading frame 166 (C20orf166), mRNA NM_080606 Homo sapiens chromosome 20 open reading frame 166 (C20orf166), mRNA NM_080606 Homo sapiens chromosome 20 open reading frame 166 (C20orf166), m	NM_080662	Homo sapiens similar to RIKEN aDNA 1810020711
MM_080659 Homo sapiens similar to RIKEN cDNA 2310030G06 gene (MGC14289), mRNA NM_080657 Homo sapiens vipirin (cig5), mRNA NM_080655 Homo sapiens similar to RIKEN cDNA 5730528L13 gene (MGC14839), mRNA NM_080654 Homo sapiens similar to RIKEN cDNA 5730528L13 gene (MGC17337), mRNA NM_080654 Homo sapiens NY-REN-41 antigen (NY-REN-41), mRNA NM_080655 Homo sapiens similar to RIKEN cDNA 4930500C14 gene (MGC9341), mRNA NM_080652 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA NM_080654 Homo sapiens regulator of G-protein signalling 6 (RGS6), mRNA NM_080655 Homo sapiens FabG (beta-ketoacyl-[acyl-carrier-protein] reductase, E coli) like (E. coli) (FABGL), mRNA NM_080626 Homo sapiens gemin 6 (GEMIN6), mRNA NM_080626 Homo sapiens chromosome 20 open reading frame 160 (C20orf160), mRNA NM_080616 Homo sapiens chromosome 20 open reading frame 112 (C20orf112), mRNA NM_080617 Homo sapiens chromosome 20 open reading frame 102 (C20orf102), mRNA NM_080603 Homo sapiens chromosome 20 open reading frame 102 (C20orf102), mRNA NM_080603 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA NM_080615 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA NM_080603 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA NM_080604 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA NM_080605 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA NM_080604 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA NM_080605 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA NM_080606 Homo sapiens chromosome 20 open reading frame 164 (C20orf164), mRNA NM_080606 Homo sapiens chromosome 20 open reading frame 165 (C20orf165), mRNA NM_080606 Homo sapiens chromosome 20 open reading frame 165 (C20orf166), mRNA NM_080606 Homo sapiens chromosome 20 open reading frame 166 (C20orf166), mRNA NM_080606 Homo sapiens chromosome 20 open reading frame 166 (C20orf166), mRNA NM_080606 Homo sapiens chromosome 20 open reading frame 166 (C20orf166), m	NM_080660	Homo sapiens similar to PIKEN aDNA 120001 1216 gene (MGC4281), mRNA
NM_080657 Homo sapiens vipirin (cig5), mRNA  NM_080655 Homo sapiens similar to RIKEN cDNA 5730528L13 gene (MGC17337), mRNA  NM_080654 Homo sapiens similar to RIKEN cDNA 5730528L13 gene (MGC17337), mRNA  NM_080653 Homo sapiens similar to RIKEN cDNA 4930500C14 gene (MGC9341), mRNA  NM_080654 Homo sapiens similar to RIKEN cDNA 4930500C14 gene (MGC9341), mRNA  NM_080655 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA  NM_080656 Homo sapiens regulator of G-protein signalling 6 (RGS6), mRNA  Homo sapiens FabG (beta-ketoacyl-[acyl-carrier-protein] reductase, E coli) like  (E. coli) (FABGL), mRNA  Homo sapiens gemin 6 (GEMIN6), mRNA  NM_080626 Homo sapiens gemin 6 (GEMIN6), mRNA  NM_080625 Homo sapiens chromosome 20 open reading frame 160 (C20orf160), mRNA  NM_080616 Homo sapiens chromosome 20 open reading frame 112 (C20orf112), mRNA  NM_080617 Homo sapiens chromosome 20 open reading frame 102 (C20orf102), mRNA  NM_080603 Homo sapiens chromosome 20 open reading frame 162 (C20orf102), mRNA  NM_080603 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA  NM_080604 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA  NM_080605 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA  NM_080604 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA  NM_080605 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA  NM_080606 Homo sapiens chromosome 20 open reading frame 164 (C20orf162), mRNA  NM_080606 Homo sapiens chromosome 20 open reading frame 164 (C20orf162), mRNA  NM_080606 Homo sapiens chromosome 20 open reading frame 165 (C20orf162), mRNA  NM_080606 Homo sapiens chromosome 20 open reading frame 165 (C20orf162), mRNA  NM_0806 Homo sapiens chromosome 20 open reading frame 165 (C20orf162), mRNA  NM_0806 Homo sapiens chromosome 20 open reading frame 165 (C20orf162), mRNA  NM_0806 Homo sapiens chromosome 20 open reading frame 165 (C20orf162), mRNA	NM_080659	
NM 080655 Homo sapiens similar to RIKEN cDNA 5730528L13 gene (MGC17337), mRNA NM 080654 Homo sapiens NY-REN-41 antigen (NY-REN-41), mRNA NM 080653 Homo sapiens similar to RIKEN cDNA 4930500C14 gene (MGC9341), mRNA NM 080652 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA NM 004296 Homo sapiens regulator of G-protein signalling 6 (RGS6), mRNA NM 014234 Homo sapiens FabG (beta-ketoacyl-[acyl-carrier-protein] reductase, E coli) like (E. coli) (FABGL), mRNA NM 080626 Homo sapiens gemin 6 (GEMIN6), mRNA NM 080625 Homo sapiens BRI3 binding protein (BRI3BP), mRNA NM 080626 Homo sapiens chromosome 20 open reading frame 160 (C20orf160), mRNA NM 080612 Homo sapiens chromosome 20 open reading frame 112 (C20orf112), mRNA NM 080607 Homo sapiens chromosome 20 open reading frame 102 (C20orf102), mRNA NM 080603 Homo sapiens chromosome 20 open reading frame 162 (C20orf102), mRNA NM 030603 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA NM 030815 Homo sapiens chromosome 20 open reading frame 162 (C20orf126), mRNA NM 030815 Homo sapiens chromosome 20 open reading frame 126 (C20orf126), mRNA NM 020841 Homo sapiens chromosome 20 open reading frame 126 (C20orf126), mRNA NM 020841 Homo sapiens chromosome 20 open reading frame 126 (C20orf126), mRNA NM 020841 Homo sapiens chromosome 20 open reading frame 126 (C20orf126), mRNA NM 020841 Homo sapiens chromosome 20 open reading frame 126 (C20orf126), mRNA		The same of the contract of th
NM 080654 Homo sapiens NY-REN-41 antigen (NY-REN-41), mRNA NM 080653 Homo sapiens similar to RIKEN cDNA 4930500C14 gene (MGC9341), mRNA NM 080652 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA NM 004296 Homo sapiens regulator of G-protein signalling 6 (RGS6), mRNA NM 014234 Homo sapiens FabG (beta-ketoacyl-[acyl-carrier-protein] reductase, E coli) like (E. coli) (FABGL), mRNA NM 080626 Homo sapiens gemin 6 (GEMIN6), mRNA NM 080625 Homo sapiens BRI3 binding protein (BRI3BP), mRNA NM 080616 Homo sapiens chromosome 20 open reading frame 160 (C20orf160), mRNA NM 080612 Homo sapiens DOS/Gab family member 3 (GAB3), mRNA NM 080603 Homo sapiens chromosome 20 open reading frame 102 (C20orf102), mRNA NM 080603 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA NM 032019 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA NM 030815 Homo sapiens chromosome 20 open reading frame 126 (C20orf126), mRNA NM 020841 Homo sapiens oxysterol binding protein-like 8 (OSBPL8), mRNA		
NM 080653 Homo sapiens similar to RIKEN cDNA 4930500C14 gene (MGC9341), mRNA NM 080652 Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA NM 004296 Homo sapiens regulator of G-protein signalling 6 (RGS6), mRNA NM 014234 Homo sapiens FabG (beta-ketoacyl-[acyl-carrier-protein] reductase, E coli) like (E. coli) (FABGL), mRNA NM 080626 Homo sapiens gemin 6 (GEMIN6), mRNA NM 080625 Homo sapiens BRI3 binding protein (BRI3BP), mRNA NM 080616 Homo sapiens chromosome 20 open reading frame 160 (C20orf160), mRNA NM 080612 Homo sapiens DOS/Gab family member 3 (GAB3), mRNA NM 080603 Homo sapiens chromosome 20 open reading frame 102 (C20orf102), mRNA NM 080603 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA NM 030815 Homo sapiens histone deacetylase 10 (HDAC10), mRNA NM 030815 Homo sapiens oxysterol binding protein-like 8 (OSBPL8), mRNA NM 020764 Homo sapiens cask-interacting protein 1 (CASERDI)		Homo sepiens Shiffiar to RIKEN cDNA 5730528L13 gene (MGC17337), mRNA
Homo sapiens similar to RIKEN cDNA 4930500C14 gene (MGC9341), mRNA NM_004296 Homo sapiens regulator of G-protein signalling 6 (RGS6), mRNA NM_014234 Homo sapiens FabG (beta-ketoacyl-[acyl-carrier-protein] reductase, E coli) like (E. coli) (FABGL), mRNA NM_080626 Homo sapiens gemin 6 (GEMIN6), mRNA NM_080625 Homo sapiens bRI3 binding protein (BRI3BP), mRNA NM_080616 Homo sapiens chromosome 20 open reading frame 160 (C20orf160), mRNA NM_080612 Homo sapiens DOS/Gab family member 3 (GAB3), mRNA NM_080607 Homo sapiens chromosome 20 open reading frame 102 (C20orf102), mRNA NM_080603 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA NM_080603 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA NM_080603 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA NM_080603 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA NM_080604 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA NM_080604 Homo sapiens chromosome 20 open reading frame 126 (C20orf162), mRNA NM_080605 Homo sapiens chromosome 20 open reading frame 126 (C20orf162), mRNA NM_080604 Homo sapiens chromosome 20 open reading frame 126 (C20orf126), mRNA NM_080605 Homo sapiens chromosome 20 open reading frame 126 (C20orf126), mRNA NM_080606 Homo sapiens chromosome 20 open reading frame 126 (C20orf126), mRNA NM_080606 Homo sapiens chromosome 20 open reading frame 126 (C20orf126), mRNA NM_080606 Homo sapiens chromosome 20 open reading frame 126 (C20orf126), mRNA NM_080606 Homo sapiens chromosome 20 open reading frame 126 (C20orf126), mRNA NM_080606 Homo sapiens chromosome 20 open reading frame 126 (C20orf126), mRNA NM_080606 Homo sapiens chromosome 20 open reading frame 126 (C20orf126), mRNA	NM 080653	
Homo sapiens regulator of G-protein signalling 6 (RGS6), mRNA Homo sapiens FabG (beta-ketoacyl-[acyl-carrier-protein] reductase, E coli) like (E. coli) (FABGL), mRNA Homo sapiens gemin 6 (GEMIN6), mRNA Homo sapiens BRI3 binding protein (BRI3BP), mRNA Homo sapiens chromosome 20 open reading frame 160 (C20orf160), mRNA Homo sapiens chromosome 20 open reading frame 112 (C20orf112), mRNA Homo sapiens DOS/Gab family member 3 (GAB3), mRNA Homo sapiens chromosome 20 open reading frame 102 (C20orf102), mRNA Homo sapiens chromosome 20 open reading frame 162 (C20orf102), mRNA Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA Homo sapiens histone deacetylase 10 (HDAC10), mRNA Homo sapiens chromosome 20 open reading frame 126 (C20orf162), mRNA Homo sapiens chromosome 20 open reading frame 126 (C20orf162), mRNA Homo sapiens chromosome 20 open reading frame 126 (C20orf162), mRNA Homo sapiens chromosome 20 open reading frame 126 (C20orf164), mRNA Homo sapiens chromosome 20 open reading frame 126 (C20orf164), mRNA Homo sapiens chromosome 20 open reading frame 126 (C20orf165), mRNA Homo sapiens cask-interacting protein-like 8 (OSBPL8), mRNA		TIVINO SADICIIS SIMILAT TO DIE EN ADMIA 4020 500 CC.
Homo sapiens FabG (beta-ketoacyl-[acyl-carrier-protein] reductase, E coli) like  (E. coli) (FABGL), mRNA  Homo sapiens gemin 6 (GEMIN6), mRNA  Homo sapiens BRI3 binding protein (BRI3BP), mRNA  Homo sapiens chromosome 20 open reading frame 160 (C20orf160), mRNA  Homo sapiens chromosome 20 open reading frame 112 (C20orf112), mRNA  Homo sapiens DOS/Gab family member 3 (GAB3), mRNA  Homo sapiens chromosome 20 open reading frame 102 (C20orf102), mRNA  MM 080603  Homo sapiens chromosome 20 open reading frame 102 (C20orf102), mRNA  NM 032019  Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA  Homo sapiens histone deacetylase 10 (HDAC10), mRNA  Homo sapiens chromosome 20 open reading frame 126 (C20orf126), mRNA  Homo sapiens chromosome 20 open reading frame 126 (C20orf126), mRNA  Homo sapiens chromosome 20 open reading frame 126 (C20orf126), mRNA  Homo sapiens chromosome 20 open reading frame 126 (C20orf126), mRNA  Homo sapiens chromosome 20 open reading frame 126 (C20orf126), mRNA  Homo sapiens cask-interacting protein l (CASKINI)  MM 020764  Homo sapiens cask-interacting protein l (CASKINI)		The supression of the first of the supression of
(E. coli) (FABGL), mRNA  NM 024775 Homo sapiens gemin 6 (GEMIN6), mRNA  NM 080626 Homo sapiens BRI3 binding protein (BRI3BP), mRNA  NM 080625 Homo sapiens chromosome 20 open reading frame 160 (C20orf160), mRNA  NM 080616 Homo sapiens chromosome 20 open reading frame 112 (C20orf112), mRNA  NM 080612 Homo sapiens DOS/Gab family member 3 (GAB3), mRNA  NM 080607 Homo sapiens chromosome 20 open reading frame 102 (C20orf102), mRNA  NM 080603 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA  NM 032019 Homo sapiens histone deacetylase 10 (HDAC10), mRNA  NM 030815 Homo sapiens chromosome 20 open reading frame 126 (C20orf162), mRNA  NM 030841 Homo sapiens chromosome 20 open reading frame 126 (C20orf126), mRNA  NM 020841 Homo sapiens oxysterol binding protein-like 8 (OSBPL8), mRNA		Homo sapiens regulator of G-protein signalling 6 (RGS6), mRNA
NM 024775 Homo sapiens gemin 6 (GEMIN6), mRNA  NM 080626 Homo sapiens BRI3 binding protein (BRI3BP), mRNA  NM 080625 Homo sapiens chromosome 20 open reading frame 160 (C20orf160), mRNA  NM 080616 Homo sapiens chromosome 20 open reading frame 112 (C20orf112), mRNA  NM 080607 Homo sapiens DOS/Gab family member 3 (GAB3), mRNA  NM 080603 Homo sapiens chromosome 20 open reading frame 102 (C20orf102), mRNA  NM 032019 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA  NM 030815 Homo sapiens chromosome 20 open reading frame 126 (C20orf162), mRNA  NM 030841 Homo sapiens chromosome 20 open reading frame 126 (C20orf126), mRNA  NM 020764 Homo sapiens oxysterol binding protein-like 8 (OSBPL8), mRNA	11111_014234	Table Sapiens rady (Dela-Ketoacyl-lacyl-carrier protein) - dela-ketoacyl-lacyl-carrier protein
NM 080626 Homo sapiens BRI3 binding protein (BRI3BP), mRNA  NM 080625 Homo sapiens chromosome 20 open reading frame 160 (C20orf160), mRNA  NM 080616 Homo sapiens chromosome 20 open reading frame 112 (C20orf112), mRNA  NM 080607 Homo sapiens DOS/Gab family member 3 (GAB3), mRNA  NM 080603 Homo sapiens chromosome 20 open reading frame 102 (C20orf102), mRNA  NM 030815 Homo sapiens histone deacetylase 10 (HDAC10), mRNA  NM 030815 Homo sapiens chromosome 20 open reading frame 126 (C20orf126), mRNA  NM 030841 Homo sapiens chromosome 20 open reading frame 126 (C20orf126), mRNA  NM 020764 Homo sapiens oxysterol binding protein-like 8 (OSBPL8), mRNA	NTM 00 4775	
Homo sapiens BRI3 binding protein (BRI3BP), mRNA  Homo sapiens chromosome 20 open reading frame 160 (C20orf160), mRNA  Homo sapiens chromosome 20 open reading frame 112 (C20orf112), mRNA  Homo sapiens DOS/Gab family member 3 (GAB3), mRNA  Homo sapiens chromosome 20 open reading frame 102 (C20orf102), mRNA  Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA  Homo sapiens histone deacetylase 10 (HDAC10), mRNA  Homo sapiens chromosome 20 open reading frame 126 (C20orf162), mRNA  Homo sapiens chromosome 20 open reading frame 126 (C20orf162), mRNA  Homo sapiens chromosome 20 open reading frame 126 (C20orf162), mRNA  Homo sapiens chromosome 20 open reading frame 126 (C20orf126), mRNA  Homo sapiens oxysterol binding protein-like 8 (OSBPL8), mRNA		Homo sapiens gemin 6 (GEMIN6), mRNA
NM 080616 Nm 080616 Homo sapiens chromosome 20 open reading frame 160 (C20orf160), mRNA NM 080612 Homo sapiens DOS/Gab family member 3 (GAB3), mRNA NM 080607 Homo sapiens chromosome 20 open reading frame 102 (C20orf102), mRNA NM 080603 Homo sapiens chromosome 20 open reading frame 102 (C20orf102), mRNA NM 032019 Homo sapiens histone deacetylase 10 (HDAC10), mRNA NM 030815 Homo sapiens chromosome 20 open reading frame 126 (C20orf162), mRNA NM 020841 Homo sapiens oxysterol binding protein-like 8 (OSBPL8), mRNA NM 020764 Homo sapiens cask-interacting protein 1 (CASENDIA)		Homo sapiens BRI3 binding protein (RRI3DD) DNA
NM 080612 Homo sapiens DOS/Gab family member 3 (GAB3), mRNA  NM 080607 Homo sapiens chromosome 20 open reading frame 102 (C20orf102), mRNA  NM 080603 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA  NM 032019 Homo sapiens histone deacetylase 10 (HDAC10), mRNA  NM 030815 Homo sapiens chromosome 20 open reading frame 126 (C20orf126), mRNA  NM 020841 Homo sapiens oxysterol binding protein-like 8 (OSBPL8), mRNA  NM 020764 Homo sapiens cask-interacting protein 1 (CASKINI)		Tiolio sapiens chromosome 20 open reading from 160 (620)
NM 080607 Homo sapiens chromosome 20 open reading frame 102 (C20orf102), mRNA NM 080603 Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA NM 032019 Homo sapiens histone deacetylase 10 (HDAC10), mRNA NM 030815 Homo sapiens chromosome 20 open reading frame 126 (C20orf126), mRNA NM 020841 Homo sapiens oxysterol binding protein-like 8 (OSBPL8), mRNA NM 020764 Homo sapiens cask-interacting protein 1 (CASKINI)		Homo sapiens chromosome 20 open reading frame 110 (C200r1160), mRNA
NM 080603 Homo sapiens chromosome 20 open reading frame 102 (C20orf102), mRNA  NM 032019 Homo sapiens histone deacetylase 10 (HDAC10), mRNA  NM 030815 Homo sapiens chromosome 20 open reading frame 126 (C20orf126), mRNA  NM 020841 Homo sapiens oxysterol binding protein-like 8 (OSBPL8), mRNA  NM 020764 Homo sapiens cask-interacting protein 1 (CASKINI)		Supromo DOD/Gab tatility themper 4 112 x 1371 toxtx
NM 032019 Homo sapiens histone deacetylase 10 (HDAC10), mRNA  NM 030815 Homo sapiens chromosome 20 open reading frame 126 (C20orf126), mRNA  NM 020841 Homo sapiens oxysterol binding protein-like 8 (OSBPL8), mRNA  NM 020764 Homo sapiens cask-interacting protein 1 (CASENDIA)	VM_080607	Homo sapiens chromosome 20 open road:
NM 032019 Homo sapiens histone deacetylase 10 (HDAC10), mRNA  NM 030815 Homo sapiens chromosome 20 open reading frame 126 (C20orf126), mRNA  NM 020841 Homo sapiens oxysterol binding protein-like 8 (OSBPL8), mRNA  NM 020764 Homo sapiens cask-interacting protein 1 (CASENDIA)	VM_080603	Homo saniens chromosome 20 open reading frame 102 (C20orf102), mRNA
MM 030815 Homo sapiens chromosome 20 open reading frame 126 (C20orf126), mRNA  MM 020841 Homo sapiens oxysterol binding protein-like 8 (OSBPL8), mRNA  MM 020764 Homo sapiens cask-interacting protein 1 (CASKINI)		- The supremental of the supreme
MM 020841 Homo sapiens enromosome 20 open reading frame 126 (C20orf126), mRNA MM 020764 Homo sapiens cask-interacting protein 1 (CASKINI)		
M 020764 Homo sapiens cask-interacting protein 1 (CASKINI) PNA	1212 02 00 13	from sapiens chromosome 20 open reading from 126 (COO)
120 Mo Sapiciis Cask-injeracting protein 1 (CASVIDIA)		
1010130   nomo sapiens chromosome 20 open reading frame 104 (C20orf104) mDNIA		210 no sapiens cask-interacting protein 1 (CACIADIA)
	111_010430	riomo sapiens chromosome 20 open reading frame 104 (C20orf104) mpara

NM 022104	Homo sapiens chromosome 20 open reading frame 67 (C20orf67), mRNA
NM 080546	Homo sapiens CDw92 antigen (CDW92), mRNA
NM 015511	Homo sapiens chromosome 20 open reading frame 4 (C20orf4), mRNA
NM 002116	Homo sapiens major histocompatibility complex, class I, A (HLA-A), mRNA
NM 023017	Homo sapiens phosphoinositide 3-kinase enhancer (PIKE), mRNA
NM 020933	Homo sapiens zinc finger protein 317 (ZNF317), mRNA
NM 005037	Homo sapiens peroxisome proliferative activated receptor, gamma (PPARG),
_	mRNA
NM 018206	Homo sapiens vacuolar protein sorting 35 (yeast) (VPS35), mRNA
NM 014003	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 38 (DDX38),
	mRNA
NM_006445	Homo sapiens PRP8 pre-mRNA processing factor 8 homolog (yeast) (PRPF8),
_	mRNA
NM 003675	Homo sapiens pre-mRNA processing factor 18 (PRP18), mRNA
NM 006214	Homo sapiens phytanoyl-CoA hydroxylase (Refsum disease) (PHYH), mRNA
NM 004374	Homo sapiens cytochrome c oxidase subunit VIc (COX6C), nuclear gene
	encoding mitochondrial protein, mRNA
NM_001863	Homo sapiens cytochrome c oxidase subunit VIb (COX6B), nuclear gene
	encoding mitochondrial protein, mRNA
NM_005205	Homo sapiens cytochrome c oxidase subunit VIa polypeptide 2 (COX6A2),
	nuclear gene encoding mitochondrial protein, mRNA
NM_004373	Homo sapiens cytochrome c oxidase subunit VIa polypeptide 1 (COX6A1),
	nuclear gene encoding mitochondrial protein, mRNA
NM_032609	Homo sapiens cytochrome c oxidase subunit IV isoform 2 (COX4I2), nuclear
	gene encoding mitochondrial protein, mRNA
NM_032489	Homo sapiens acrosin binding protein (ACRBP), mRNA
NM_080476	Homo sapiens CDC91 cell division cycle 91-like 1 (S. cerevisiae) (CDC91L1),
	mRNA
NM_080473	Homo sapiens GATA binding protein 5 (GATA5), mRNA
NM_002121	Homo sapiens major histocompatibility complex, class II, DP beta 1 (HLA-DPB1), mRNA
NM_078470	Homo sapiens COX15 homolog, cytochrome c oxidase assembly protein (yeast)
_	(COX15), nuclear gene encoding mitochondrial protein, transcript variant 1,
	mRNA
NM_004375	Homo sapiens COX11 homolog, cytochrome c oxidase assembly protein (yeast)
	(COX11), nuclear gene encoding mitochondrial protein, mRNA
NM_001303	Homo sapiens COX10 homolog, cytochrome c oxidase assembly protein, heme
	A/farnesyltransferase (yeast) (COX10), nuclear gene encoding mitochondrial
	protein, mRNA
NM_054028	Homo sapiens acyl-malonyl condensing enzyme (AMAC), mRNA
NM_032485	Homo sapiens chromosome 20 open reading frame 154 (C20orf154), mRNA
NM_033342	Homo sapiens tripartite motif-containing 7 (TRIM7), mRNA
NM_033421	Homo sapiens chromosome 20 open reading frame 161 (C20orf161), mRNA
NM_033197	Homo sapiens chromosome 20 open reading frame 114 (C20orf114), mRNA
NM_020866	Homo sapiens kelch-like 1 (Drosophila) (KLHL1), mRNA
NM_032883	Homo sapiens chromosome 20 open reading frame 100 (C20orf100), mRNA
NM_032523	Homo sapiens oxysterol binding protein-like 6 (OSBPL6), mRNA
NM_020896	Homo sapiens oxysterol binding protein-like 5 (OSBPL5), mRNA
NM_015550	Homo sapiens oxysterol binding protein-like 3 (OSBPL3), mRNA
NM_031473	Homo sapiens carnitine deficiency-associated gene expressed in ventricle 1
	(CDV-1), mRNA
NM_030801	Homo sapiens MAGE-E1 protein (MAGE-E1), mRNA

NM_025128	Homo sapiens MUS81 endonuclease (MUS81), mRNA
NM_024958	Homo sapiens chromosome 20 open reading frame 98 (C20orf98) mPNA
NM_024663	Homo sapiens aminopeptidase-like 1 (NPEPL1) mRNA
NM_024586	Homo sapiens oxysterol binding protein-like 9 (OSBPI 9) mDNA
NM_024120	Homo sapiens chromosome 20 open reading frame 7 (C20orf7), mPNIA
NM_022776	Homo sapiens oxysterol binding protein-like 11 (OSRPI 11) mpN/A
NM_022109	Hollo Sapiens CDw92 antigen (CDW92) mRNA
NM_022088	Homo sapiens zinc finger protein 338 (ZNF338) mRNA
NM_021158	Homo sapiens chromosome 20 open reading frame 97 (C20orf07) mDNIA
NM_021232	110110 sapiens profine denydrogenase (oxidase) 2 (PRODH2) mDNA
NM_021220	Hollo sapiens zinc finger protein 339 (ZNF339) mRNA
NM_021039	Homo sapiens S100 calcium binding protein A14 (calgizzarin) (S100A14),
	IIICIVA
NM_020659	Homo sapiens tweety homolog 1 (Drosophila) (TTYH1), mRNA
NM_018972	Homo sapiens ganglioside-induced differentiation-associated protein 1
	(ODAFI), MKNA
NM_017921	Homo sapiens hypothetical protein FLI20657 (NPL4) mRNA
NM_017784	Homo sapiens oxysterol binding protein-like 10 (OSBPI 10) mPNIA
NM_017731	Homo sapiens oxysterol binding protein-like 7 (OSRPI 7) mPNIA
NM_018209	Homo sapiens ADP-ribosylation factor 1 GTPase activating protein
	(AKTIGAP), MKNA
NM_018102	Homo sapiens zinc finger protein 334 (ZNF334) mPNA
NM_015891	Homo sapiens pre-mRNA splicing factor 17 (PRP17), mRNA
NM_016599	Homo sapiens myozenin 2 (MYOZ2), mRNA
NM_014962	Homo sapiens BTB (POZ) domain containing 3 (RTRD3) mPNA
NM_014835	Homo sapiens oxysterol binding protein-like 2 (OSBPL2), mRNA
NM_014723	Homo sapiens syntaphilin (SNPH), mRNA
NM_014183	Homo sapiens dynein light chain 2A (DNLC2A), mRNA
NM_014055	Homo sapiens carnitine deficiency-associated gene expressed in ventricle 1
	(CDV-1), IIKNA
NM_014477	Homo sapiens chromosome 20 open reading frame 10 (C20orf10), mRNA
NM_012261	Tiomo sapiens chromosome 20 open reading frame 103 (C20orf102) DNA
NM_013369	Homo sapiens DNA (cytosine-5-)-methyltransferase 3-like (DNMT3L), mRNA
NM_012469	Tromo sapiens chromosome 20 open reading frame 14 (C20orf14) mDNA
NM_012291	Hollo sapiens extra spindle poles like 1 (S cerevisiae) (FSPI 1) mpNIA
NM_007002	fromo sapiens agnesion regulating molecule 1 (ADRM1) mDNA
NM_006809	Homo sapiens translocase of outer mitochondrial membrane 34 (TOMM34),
	IIIIIIVA
NM_006813	Homo sapiens proline rich 2 (PROL2), mRNA
NM_002509	Homo sapiens NK2 transcription factor homolog B (Drosophila) (NKV2B)
	IIICIVA
NM_080474	Homo sapiens serine (or cysteine) proteinase inhibitor, clade B (ovalbumin),
	member 12 (SERPINB12), mRNA
NM_006009	Homo sapiens tubulin, alpha 3 (TUBA3), mRNA
NM_003463	Homo sapiens protein tyrosine phosphatase type IVA member 1 (PTPAA1)
	MICHA
NM_019888	Homo sapiens melanocortin 3 receptor (MC3R), mRNA
NM_001846	Homo sapiens collagen, type IV, alpha 2 (COI 4A2) mRNA:
NM_079422	Homo sapiens myosin, light polypeptide 1, alkali; skeletal, fast (MYL1),
	transcript variant 31, mrnA
NM_079420	Homo sapiens myosin, light polypeptide 1, alkali; skeletal, fast (MYL1),
	transcript variant 1f, mRNA

NM_000795	Homo sapiens dopamine receptor D2 (DRD2), transcript variant 1, mRNA
NM_016574	Homo sapiens dopamine receptor D2 (DRD2), transcript variant 2, mRNA
NM_079837	Homo sapiens BTG3 associated nuclear protein (BANP), transcript variant 2, mRNA
NM_017869	Homo sapiens BTG3 associated nuclear protein (BANP), transcript variant 1, mRNA
NM_079425	Homo sapiens myosin, light polypeptide 6, alkali, smooth muscle and non-muscle (MYL6), transcript variant 3, mRNA
NM_079424	Homo sapiens myosin, light polypeptide 6, alkali, smooth muscle and non-muscle (MYL6), transcript variant 4, mRNA
NM_079423	Homo sapiens myosin, light polypeptide 6, alkali, smooth muscle and non-muscle (MYL6), transcript variant 2, mRNA
NM_021019	Homo sapiens myosin, light polypeptide 6, alkali, smooth muscle and non-muscle (MYL6), transcript variant 1, mRNA
NM 004509	Homo sapiens SP110 nuclear body protein (SP110), transcript variant a, mRNA
NM 080424	Homo sapiens SP110 nuclear body protein (SP110), transcript variant c, mRNA
NM 004510	Homo sapiens SP110 nuclear body protein (SP110), transcript variant b, mRNA
NM 004574	Homo sapiens peanut-like 2 (Drosophila) (PNUTL2), transcript variant 1, mRNA
NM 080417	Homo sapiens peanut-like 2 (Drosophila) (PNUTL2), transcript variant 4, mRNA
NM_080416	Homo sapiens peanut-like 2 (Drosophila) (PNUTL2), transcript variant 3, mRNA
NM 080415	Homo sapiens peanut-like 2 (Drosophila) (PNUTL2), transcript variant 2, mRNA
NM 002117	Homo sapiens major histocompatibility complex, class I, C (HLA-C), mRNA
NM 005514	Homo sapiens major histocompatibility complex, class I, B (HLA-B), mRNA
NC_001807	Homo sapiens mitochondrion, complete genome
NM 080489	Homo sapiens syndecan binding protein (syntenin) 2 (SDCBP2), mRNA
NM 001997	Homo sapiens Finkel-Biskis-Reilly murine sarcoma virus (FBR-MuSV)
11111_001557	ubiquitously expressed (fox derived); ribosomal protein S30 (FAU), mRNA
NM 057179	Homo sapiens likely ortholog of mouse and rat twist-related bHLH protein
1111_037173	Dermo-1 (DERMO1), mRNA
NM 001008	Homo sapiens ribosomal protein S4, Y-linked (RPS4Y), mRNA
NM 001007	Homo sapiens ribosomal protein S4, X-linked (RPS4X), mRNA
NM 005192	Homo sapiens cyclin-dependent kinase inhibitor 3 (CDK2-associated dual
<u> </u>	specificity phosphatase) (CDKN3), mRNA
NM_079421	Homo sapiens cyclin-dependent kinase inhibitor 2D (p19, inhibits CDK4) (CDKN2D), transcript variant 2, mRNA
NM_001800	Homo sapiens cyclin-dependent kinase inhibitor 2D (p19, inhibits CDK4) (CDKN2D), transcript variant 1, mRNA
NM_078626	Homo sapiens cyclin-dependent kinase inhibitor 2C (p18, inhibits CDK4) (CDKN2C), transcript variant 2, mRNA
NM_001262	Homo sapiens cyclin-dependent kinase inhibitor 2C (p18, inhibits CDK4) (CDKN2C), transcript variant 1, mRNA
NM_078487	Homo sapiens cyclin-dependent kinase inhibitor 2B (p15, inhibits CDK4) (CDKN2B), transcript variant 2, mRNA
NM_004936	Homo sapiens cyclin-dependent kinase inhibitor 2B (p15, inhibits CDK4) (CDKN2B), transcript variant 1, mRNA
NM 004896	Homo sapiens vacuolar protein sorting 26 (yeast) (VPS26), mRNA
NM 052945	Homo sapiens BAFF receptor (BAFFR), mRNA
NM 022648	Homo sapiens tensin (TNS), mRNA
1 1111 022070	
NM_078480	Homo sapiens fuse-binding protein-interacting repressor (SIAHBP1), transcript variant 1, mRNA

NM_004740	Homo sapiens TGFB1-induced anti-
	2. mRNA 251 B1-induced anti-apoptotic factor 1 (TIAF1), transcript variation
NM_078471	Homo sapiens TGFR1 induced
	Homo sapiens TGFB1-induced anti-apoptotic factor 1 (TIAF1), transcript varia
NM 001852	Homo saniens colleges 4
NM_078485	The supreme su
NM_001851	The suprems confident type IX alpha 1 (COT 0.4.1)
NM_054026	Homo sapiens collagen, type IX, alpha 1 (COL9A1), transcript variant 2, mRNA Homo sapiens CCR4-NOT transcription accounts.
	Homo sapiens CCR4-NOT transcription complex, subunit 7 (CNOT7), transcript variant 2, mRNA
NM_013354	Homo series COD (1997), transcrip
	Homo sapiens CCR4-NOT transcription complex, subunit 7 (CNOT7), transcription variant 1, mRNA
NM_004064	variant 1, mRNA  Home complex, subunit 7 (CNOT7), transcrip
	Homo sapiens cyclin-dependent kinase inhibitor 1B (p27, Kip1) (CDKN1B), mRNA
NM_000389	Home or
- 1212_000303	Homo sapiens cyclin-dependent kinase inhibitor 1A (p21, Cip1) (CDKN1A), transcript variant 1, mRNA
NM_078467	transcript variant 1, mRNA
1111_076407	Homo sapiens cyclin-dependent kinase inhibitor 1A (p21, Cip1) (CDKN1A), transcript variant 2, mRNA
NM_003936	transcript variant 2, mRNA
14141_003930	Homo sapiens cyclin-dependent kinase 5, regulatory subunit 2 (p39) (CDK5R2), mRNA
NM_004642	mRNA (DS9) (CDK5R2),
	Homo sapiens CDK2-associated protein 1 (CDK2AP1), mRNA
NM_078481	
NM_001784	
NM_080432	Homo sapiens vacuolar protein sorting protein 18 (VPS18), transcript variant 2, mRNA mRNA
ND ( 0000	mRNA protein 18 (VPS18), transcript variant 2,
NM_020857	Homo sapiens vacuolar protein sorting protein 18 (VPS18), transcript variant 1, mRNA
NT 6 000 11	mRNA protein sorting protein 18 (VPS18), transcript variant 1,
NM_080414	Homo sapiens vacuolar protein sorting 16 (yeast) (VPS16), transcript variant 2, mRNA
The opposite	mRNA (VPS16), transcript variant 2,
NM_080413	Homo sapiens vacuolar protein sorting 16 (yeast) (VPS16), transcript variant 3, mRNA
Th. 6.000	mRNA protein softing to (yeast) (VPS16), transcript variant 3,
NM_022575	Homo sapiens vacuolar protein sorting 16 (yeast) (VPS16), transcript variant 1, mRNA
72.5	mRNA (yeast) (VPS16), transcript variant 1,
M_021729	Homo sapiens vacuolar protein sorting 11 (yeast) (VPS11), mRNA
M_005806	
JM_012106	Homo sapiens binder of Arl Two (BART1), mRNA
IM_006095	Homo sapiens ATPase aminopheculali il
	Homo sapiens ATPase, aminophospholipid transporter (APLT), Class I, type 8A, member 1 (ATP8A1), mRNA
M_058241	Homo sapiens cyclin T2 (CCNT2)
M_001241	Homo sapiens cyclin T2 (CCNT2), transcript variant b, mRNA Homo sapiens cyclin T2 (CCNT2)
M_001240	Homo sapiens cyclin T2 (CCNT2), transcript variant b, mRNA Homo sapiens cyclin T1 (CCNT1), mRNA Homo sapiens cyclin T1 (CCNT1), mRNA
	Homo sapiens twist homolog (acrocephalosyndactyly 3; Saethre-Chotzen syndrome) (Drosophila) (TWIST), mRNA
M_080475 ]	Homo saniens serine (or cyrtain), mKNA
	Homo sapiens serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 11 (SERPINB11), mRNA
	Homo sapiens caspase recruitment domain protein 12 (CARD12), mRNA
M_007049 F	Homo sapiens centaurin, beta 2 (CENTB2), mRNA
	Tomo sapiens outyrophilin subfamily 2 months A 1 (Dom to
M_078476 H	rariant 1, mRNA  John Sanita I. (B1N2A1), transcript
VI ()/X4I/h	tomo sapiens butyrophilin, subfamily 2 member 4.1 (DTND 4.1)
1	nmiont 2 Data
v	Iomo sapiens butyrophilin, subfamily 2, member A1 (BTN2A1), transcript Iomo sapiens EphB4 (EPHB4), mRNA

NM_004443	Homo sapiens EphB3 (EPHB3), mRNA
NM_004442	Homo sapiens EphB2 (EPHB2), transcript variant 1, mRNA
NM_017449	Homo sapiens EphB2 (EPHB2), transcript variant 2, mRNA
NM_004535	Homo sapiens myelin transcription factor 1 (MYT1), mRNA
NM_006800	Homo sapiens male-specific lethal 3-like 1 (Drosophila) (MSL3L1), transcript variant 3, mRNA
NM_078630	Homo sapiens male-specific lethal 3-like 1 (Drosophila) (MSL3L1), transcript variant 2, mRNA
NM_078629	Homo sapiens male-specific lethal 3-like 1 (Drosophila) (MSL3L1), transcript variant 1, mRNA
NM_078628	Homo sapiens male-specific lethal 3-like 1 (Drosophila) (MSL3L1), transcript variant 4, mRNA
NM_080431	Homo sapiens actin related protein M2 (ARPM2), mRNA
NM 080430	Homo sapiens selenoprotein SelM (SELM), mRNA
NM 052944	Homo sapiens putative sodium-coupled cotransporter RKST1 (RKST1), mRNA
NM_024831	Homo sapiens nuclear receptor coactivator 6 interacting protein (NCOA6IP), mRNA
NM_032803	Homo sapiens solute carrier family 7 (cationic amino acid transporter, y+system), member 3 (SLC7A3), mRNA
NM_080385	Homo sapiens carboxypeptidase A5 (CPA5), mRNA
NM_016476	Homo sapiens APC11 anaphase promoting complex subunit 11 homolog (yeast) (ANAPC11), mRNA
NM_080389	Homo sapiens defensin, beta 4 (DEFB4), mRNA
NM 032646	Homo sapiens tweety homolog 2 (Drosophila) (TTYH2), mRNA
NM 006928	Homo sapiens silver homolog (mouse) (SILV), mRNA
NM 080390	Homo sapiens my048 protein (my048), mRNA
NM 080388	Homo sapiens hypothetical protein MGC17528 (MGC17528), mRNA
NM 080387	Homo sapiens C-type lectin-like receptor (CLEC-6), mRNA
NM_080284	Homo sapiens ATP-binding cassette, sub-family A (ABC1), member 6 (ABCA6), mRNA
NM_080283	Homo sapiens ATP-binding cassette, sub-family A (ABC1), member 9 (ABCA9), mRNA
NM_080282	Homo sapiens ATP-binding cassette, sub-family A (ABC1), member 10 (ABCA10), mRNA
NM_006549	Homo sapiens calcium/calmodulin-dependent protein kinase kinase 2, beta (CAMKK2), mRNA
NM_007200	Homo sapiens A kinase (PRKA) anchor protein 13 (AKAP13), mRNA
NM_002476	Homo sapiens myosin, light polypeptide 4, alkali; atrial, embryonic (MYL4), mRNA
NM_001853	Homo sapiens collagen, type IX, alpha 3 (COL9A3), mRNA
NM_006001	Homo sapiens tubulin, alpha 2 (TUBA2), transcript variant 1, mRNA
NM_079836	Homo sapiens tubulin, alpha 2 (TUBA2), transcript variant 2, mRNA
NM_006000	Homo sapiens tubulin, alpha 1 (testis specific) (TUBA1), mRNA
NM_004376	Homo sapiens COX15 homolog, cytochrome c oxidase assembly protein (yeast) (COX15), nuclear gene encoding mitochondrial protein, transcript variant 2, mRNA
NM_024407	Homo sapiens NADH dehydrogenase (ubiquinone) Fe-S protein 7 (20kD) (NADH-coenzyme Q reductase) (NDUFS7), mRNA
NM_078625	Homo sapiens vanin 3 (VNN3), transcript variant 2, mRNA
NM 018399	Homo sapiens vanin 3 (VNN3), transcript variant 1, mRNA
NM 078488	Homo sapiens vanin 2 (VNN2), transcript variant 2, mRNA
NM 004665	Homo sapiens vanin 2 (VNN2), transcript variant 1, mRNA

NM_013245	Homo caniena vicanala
NM_058240	
	- 20 me suprens solute carrier tamily x (sodium coloium and
NM_033262	
	Homo sapiens solute carrier family 8 (sodium-calcium exchanger), member 3 (SLC8A3), transcript variant a, mRNA
NM 004869	- 1 - 2 ); stanscript variant a. mk NA
NM 078474	The supplies of the supplies o
NM_025141	The Digital Alberta Country of the Digital Alberta Country of the Digital Country of the Di
NM 078473	1 - 20 III O Supremo DDI - IIKE DIOIEIN / (RI D/) trongomint
NM 031940	1 220 Ho Supicins DDI -like Drotein ( R) Pl) transposint 1
NM_020749	- 1 - 1 - 1 - 1 - 1 - 1 - 1 - 1 - 1 - 1
NM_018672	- 1 A TTD1\ Date Did A 1 A 1 CUCUIOI - Interacting protein 1 (A TTD1\ Date
_	Homo sapiens ATP-binding cassette, sub-family A (ABC1), member 5 (ABCA5), mRNA
NM_020177	Homo saniens feminization 11.
NM_002088	Homo sapiens edutements with the Homo sapiens edutement with the Homo sapiens edutement with the Homo sapiens edutement with the Homo sapiens education of the Homo sapiens education in the Homo sapiens
NM_006835	Homo sapiens glutamate receptor, ionotropic, kainate 5 (GRIK5), mRNA  Homo sapiens cyclin I (CCNI), mRNA
NM_001239	Tarable suprems cyclin I (CCNI) mRNA
NM 014286	Homo sapiens cyclin H (CCNH), mRNA Homo sapiens frequenin homo sapiens
NM_006650	Homo sapiens frequenin homolog (Drosophila) (FREQ), mRNA
NM_006651	Homo sapiens complexin 2 (CPLX2), mRNA  Homo sapiens complexin 2 (CPLX2), mRNA
NM_006463	Homo sapiens complexin 1 (CPLX1), mRNA
	Homo sapiens associated molecule with the SH3 domain of STAM (AMSH), mRNA
NM_001850	
NM 000094	Homo sapiens collagen, type VIII, alpha 1 (COL8A1), mRNA
_	1 220110 Supicits Collayen, Type VII 2 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
NM_000077	
	Homo sapiens cyclin-dependent kinase inhibitor 2A (melanoma, p16, inhibits CDK4) (CDKN2A) transcript verient 1 PNA
NM_058197	
	Homo sapiens cyclin-dependent kinase inhibitor 2A (melanoma, p16, inhibits CDK4) (CDKN2A) transcript variant 2 PNA
NM_058196	
	Homo sapiens cyclin-dependent kinase inhibitor 2A (melanoma, p16, inhibits CDK4) (CDKN2A), transcript variant 2, mRNA
NM_058195	
	Homo sapiens cyclin-dependent kinase inhibitor 2A (melanoma, p16, inhibits CDK4) (CDKN2A), transcript variant 4, mRNA
NM_014800	
	Homo sapiens engulfment and cell motility 1 (ced-12 homolog, C. elegans) (ELMO1), mRNA
NM_079834	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
NM_019110	Homo sapiens secretory carrier membrane protein 4 (SCAMP-4), mRNA  Homo sapiens hypothetical protein Pl. 278 6 778 1
NM_022086	The suprems mypoinicitical project P1 p373c6 (P1 p27200) Part
	Homo sapiens engulfment and cell motility 2 (ced-12 homolog, C. elegans) (ELMO2), mRNA
NM_058183	
NM_003103	Homo sapiens SON DNA binding protein (SON), mRNA Homo sapiens SON DNA binding protein (SON), mRNA
NM_030767	Homo sapiens SON DNA binding protein (SON), mRNA  Homo sapiens AT-book transcription (SON), mRNA
VM_058191	Homo sapiens AT-hook transcription factor AKNA (AKNA), mRNA  Homo sapiens chromosome 21 areas at the factor AKNA (AKNA), mRNA
VM_015657	The supposed of the control of the c
	Homo sapiens ATP-binding cassette, sub-family A (ABC1), member 12 (ABCA12), mRNA
VM_020427	
VM_021638	Homo sapiens actin file and the Homo sapiens actin file actin file and the Homo sapiens actin file ac
JM_005782	Homo sapiens actin filament associated protein (AFAP), mRNA  Homo sapiens transcriptional
	The supports it all SCI ID HODRI COSC CITY of CAT TO ALL TO ALL
	270 Mo Sapiens Al AP-associated sperm protoin (ASD) - Dall
	Homo sapiens alveolar soft part sarcoma chromosome region, candidate 1 (ASPSCR1), mRNA
	Homo sapiens zinc finger protein 354B (ZNF354B), mRNA
	Tiomo sapiens zine finger protein 354B (ZNE354B) mDNA

NM_021935	Homo sapiens homolog of mouse Bv8 (Bombina variegata 8 kDa); prokineticin 2 precursor (BV8), mRNA
NM 015399	Homo sapiens breast cancer metastasis-suppressor 1 (BRMS1), mRNA
NM 007073	Homo sapiens blood vessel epicardial substance (BVES), mRNA
NM_017726	Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 14D (PPP1R14D), mRNA
NM_006451	Homo sapiens polyadenylate binding protein-interacting protein 1 (PAIP1), mRNA
NM 018073	Homo sapiens SSA protein SS-56 (SS-56), mRNA
NM 032812	Homo sapiens tumor endothelial marker 7-related precursor (TEM7R), mRNA
NM 022748	Homo sapiens tumor endothelial marker 6 (TEM6), mRNA
NM 032777	Homo sapiens tumor endothelial marker 5 precursor (TEM5), mRNA
NM_022779	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 31 (DDX31), mRNA
NM_018454	Homo sapiens nucleolar protein ANKT (ANKT), mRNA
NM_016489	Homo sapiens uridine 5' monophosphate hydrolase 1 (UMPH1), mRNA
NM_078483	Homo sapiens lysosomal amino acid transporter 1 (LYAAT1), mRNA
NM_019606	Homo sapiens hypothetical protein FLJ20257 (FLJ20257), mRNA
NM 015256	Homo sapiens fatty-acid-Coenzyme A ligase, long-chain 6 (FACL6), mRNA
NM_003393	Homo sapiens wingless-type MMTV integration site family, member 8B (WNT8B), mRNA
NM_058244	Homo sapiens wingless-type MMTV integration site family, member 8A (WNT8A), transcript variant 2, mRNA
NM_058238	Homo sapiens wingless-type MMTV integration site family, member 7B (WNT7B), mRNA
NM_004625	Homo sapiens wingless-type MMTV integration site family, member 7A (WNT7A), mRNA
NM 058242	Homo sapiens keratin 6C (KRT6C), mRNA
NM 005555	Homo sapiens keratin 6B (KRT6B), mRNA
NM 005554	Homo sapiens keratin 6A (KRT6A), mRNA
NM_058207	Homo sapiens sperm associated antigen 11 (SPAG11), transcript variant E, mRNA
NM_058206	Homo sapiens sperm associated antigen 11 (SPAG11), transcript variant B, mRNA
NM_058203	Homo sapiens sperm associated antigen 11 (SPAG11), transcript variant C, mRNA
NM_058202	Homo sapiens sperm associated antigen 11 (SPAG11), transcript variant H, mRNA
NM_058201	Homo sapiens sperm associated antigen 11 (SPAG11), transcript variant D, mRNA
NM_058200	Homo sapiens sperm associated antigen 11 (SPAG11), transcript variant G, mRNA
NM_016512	Homo sapiens sperm associated antigen 11 (SPAG11), transcript variant A, mRNA
NM_057180	Homo sapiens vacuolar protein sorting 29 (yeast) (VPS29), transcript variant 2, mRNA
NM_016226	Homo sapiens vacuolar protein sorting 29 (yeast) (VPS29), transcript variant 1, mRNA
NM_053004	Homo sapiens guanine nucleotide binding protein (G protein), beta polypeptide 1-like (GNB1L), mRNA
NM_003902	Homo sapiens far upstream element (FUSE) binding protein 1 (FUBP1), mRNA
NM 058217	Homo sapiens RAD51 homolog C (S. cerevisiae) (RAD51C), transcript variant

NIM OF COLE	3, mRNA
NM_058216	The same of the sa
ND4 000056	
NM_002876	
377.5	
NM_058179	
NM_021154	Homo sapiens phosphoserine aminotransferase (PSA), transcript variant 2, mRNA
NM_078469	Homo sapiens BRCA2 and CDKN1A interacting protein (BCCIP), transcript variant C. mRNA
NM_078468	Homo sapiens BRCA2 and CDKN1A interacting protein (BCCIP), transcript variant B, mRNA
NM_016567	Homo sapiens BRCA2 and CDKN1A interacting protein (BCCIP), transcript variant A, mRNA
	variant A, mRNA
NM_058177	
_	Homo sapiens histone deacetylase 9 (HDAC9-PENDING), transcript variant 2, mRNA
NM_058176	
	Homo sapiens histone deacetylase 9 (HDAC9-PENDING), transcript variant 1, mRNA
NM_022110	
NM_012181	Homo sapiens FK506 binding protein like (FKBPL), mRNA
NM_003602	Tionio sapiciis FN 300 hinding protein 9 (2017) (Elepho)
NM_004117	Tromo supiciis I'N 300 Dinging profein 6 (36kD) (TRODO)
	- 20 Mio Supicing Little Different Clark DDC - Data
NM_002014	1101110 Sapicits FROUD hinding protein 4 (501-D) (FRDDA)
NM_057092	220 Me suprens PK 500 unding protein 2 (13kl)) (FK PD2) two-
)TI ( 004455	
NM_004470	Homo sapiens FK506 binding protein 2 (13kD) (FKBP2), transcript variant 1, mRNA
3B 5 00 11 1	
NM_004116	Homo sapiens FK506 binding protein 1B (12.6 kD) (FKBP1B), transcript variant 1, mRNA
NM_054033	Homo sapiens FK506 binding protein 1B (12.6 kD) (FKBP1B), transcript variant 2, mRNA
NM_000801	Homo sapiens FK506 binding protein 1A (12kD) (FKBP1A), transcript variant 12B, mRNA
	12B, mRNA
NM_054014	Homo sapiens FK506 binding protein 1A (12kD) (FKBP1A), transcript variant 12A, mRNA
	12A, mRNA
NM_057175	Homo saniens hypothetical protein DV VICO to
	Homo sapiens hypothetical protein FLJ13340 (FLJ13340), transcript variant 1, mRNA
NM 025085	
	Homo sapiens hypothetical protein FLJ13340 (FLJ13340), transcript variant 2, mRNA
NM_014708	
NM_058199	Homo sapiens kinetochore associated 1 (KNTC1), mRNA
VM_014279	Tromo sapiens offactomedin   (O) FM1) transcript residue 2
VM 057174	
1171_03/1/4	220 Mo Supicits per 0x150 mai piogenesis factor 16 (DEV16)
JM 022110	
VM 033118	Homo sapiens myosin light chain kinase 2, skeletal muscle (MYLK2), mRNA
1	From the full till till till till till till till t
1	
	TT
IM_022549	nomo sapiens fasciculation and elongation proteins.
JM_022549	Homo sapiens fasciculation and elongation protein zeta 1 (zygin I) (FEZ1), transcript variant 2, mRNA  Homo sapiens WD repeat domain 1 (WDR1), transcript variant 2, mRNA

NM_017491	Homo sapiens WD repeat domain 1 (WDR1), transcript variant 1, mRNA
NM 001862	Homo sapiens cytochrome c oxidase subunit Vb (COX5B), nuclear gene
_	encoding mitochondrial protein, mRNA
NM 004255	Homo sapiens cytochrome c oxidase subunit Va (COX5A), nuclear gene
_	encoding mitochondrial protein, mRNA
NM 057162	Homo sapiens kelch-like 4 (Drosophila) (KLHL4), transcript variant 2, mRNA
NM 033427	Homo sapiens cortactin binding protein 2 (CORTBP2), mRNA
NM 001799	Homo sapiens cyclin-dependent kinase 7 (MO15 homolog, Xenopus laevis, cdk-
	activating kinase) (CDK7), mRNA
NM 057089	Homo sapiens adaptor-related protein complex 1, sigma 1 subunit (AP1S1),
_	transcript variant 2, mRNA
NM 001283	Homo sapiens adaptor-related protein complex 1, sigma 1 subunit (AP1S1),
	transcript variant 1, mRNA
NM_005148	Homo sapiens unc-119 homolog (C. elegans) (UNC119), transcript variant 1,
	mRNA
NM_054035	Homo sapiens unc-119 homolog (C. elegans) (UNC119), transcript variant 2,
_	mRNA
NM_017675	Homo sapiens protocadherin LKC (PC-LKC), mRNA
NM_002401	Homo sapiens mitogen-activated protein kinase kinase kinase 3 (MAP3K3),
	mRNA
NM_003728	Homo sapiens unc-5 homolog B (C. elegans) (UNC5C), mRNA
NM_004673	Homo sapiens angiopoietin-like 1 (ANGPTL1), mRNA
NM_054016	Homo sapiens FUS interacting protein (serine-arginine rich) 1 (FUSIP1),
_	transcript variant 2, mRNA
NM_006625	Homo sapiens FUS interacting protein (serine-arginine rich) 1 (FUSIP1),
_	transcript variant 1, mRNA
NM_054027	Homo sapiens ankylosis, progressive homolog (mouse) (ANKH), transcript
	variant 2, mRNA
NM_019847	Homo sapiens ankylosis, progressive homolog (mouse) (ANKH), transcript
	variant 1, mRNA
NM_006363	Homo sapiens Sec23 homolog B (S. cerevisiae) (SEC23B), transcript variant 1,
	mRNA
NM_032986	Homo sapiens Sec23 homolog B (S. cerevisiae) (SEC23B), transcript variant 3,
-	mRNA
NM_032985	Homo sapiens Sec23 homolog B (S. cerevisiae) (SEC23B), transcript variant 2,
	mRNA
NM_053285	Homo sapiens tektin 1 (TEKT1), mRNA
NM_018440	Homo sapiens phosphoprotein associated with glycosphingolipid-enriched
	microdomains (PAG), mRNA
NM_014479	Homo sapiens ADAM-like, decysin 1 (ADAMDEC1), mRNA
NM_016545	Homo sapiens immediate early response 5 (IER5), mRNA
NM_052820	Homo sapiens coronin, actin binding protein, 2A (CORO2A), transcript variant
37.	2, mRNA
NM_003389	Homo sapiens coronin, actin binding protein, 2A (CORO2A), transcript variant
27.6.05.55	1, mRNA
NM_032587	Homo sapiens caspase recruitment domain family, member 6 (CARD6), mRNA
NM_052814	Homo sapiens caspase recruitment domain family, member 9 (CARD9),
	transcript variant 2, mRNA
NM_052813	Homo sapiens caspase recruitment domain family, member 9 (CARD9),
	transcript variant 1, mRNA
NM_022352	Homo sapiens caspase recruitment domain family, member 9 (CARD9),
<del> </del>	transcript variant 3, mRNA

)77.6 072070	
NM_052978	Homo sapiens tripartite motif-containing 9 (TRIM9), transcript variant 2, mRNA
NM_015163	Homo sapiens tripartite motif-containing 9 (TRIM9), transcript variant 1, mRNA
NM_052840	Homo sapiens bruno-like 6, RNA binding protein (Drosophila) (BRUNOL6),
NM 000967	mRNA
NM 015125	Homo sapiens ribosomal protein L3 (RPL3), mRNA
NM 018256	Homo sapiens capicua homolog (Drosophila) (CIC), mRNA
NM_016601	Homo sapiens WD repeat domain 12 (WDR12), mRNA
14141_010001	Homo sapiens potassium channel, subfamily K, member 9 (TASK-3) (KCNK9), mRNA
NM 033415	Homo sapiens hypothetical gene MGC19595 (MGC19595), mRNA
NM_001253	Homo sapiens CDC5 cell division cycle 5-like (S. pombe) (CDC5L), mRNA
NM_007065	Homo sapiens CDC37 cell division cycle 37 homolog (S. cerevisiae) (CDC37), mRNA
NM 003504	Homo sapiens CDC45 cell division cycle 45-like (S. cerevisiae) (CDC45L),
1111_005504	mRNA
NM_006035	Homo sapiens CDC42 binding protein kinase beta (DMPK-like) (CDC42BPB), mRNA
NM_044472	Homo sapiens cell division cycle 42 (GTP binding protein, 25kD) (CDC42), transcript variant 2, mRNA
NM_001791	Homo sapiens cell division cycle 42 (GTP binding protein, 25kD) (CDC42),
77.000	transcript variant 1, mRNA
NM_001254	Homo sapiens CDC6 cell division cycle 6 homolog (S. cerevisiae) (CDC6), mRNA
NM 022894	Homo sapiens poly(A) polymerase gamma (PAPOLG), mRNA
NM_033655	Homo sapiens cell recognition molecule CASPR3 (CASPR3), transcript variant
_	1, mRNA
NM_024879	Homo sapiens cell recognition molecule CASPR3 (CASPR3), transcript variant 2, mRNA
NM_012115	Homo sapiens CASP8 associated protein 2 (CASP8AP2), mRNA
NM 012173	Homo sapiens F-box only protein 25 (FBXO25), mRNA
NM 033624	Homo sapiens F-box only protein 21 (FBXO21), transcript variant 1, mRNA
NM_015002	Homo sapiens F-box only protein 21 (FBXO21), transcript variant 2, mRNA
NM_033625	Homo sapiens ribosomal protein L34 (RPL34), transcript variant 2, mRNA
NM_000995	Homo sapiens ribosomal protein L34 (RPL34), transcript variant 1, mRNA
NM_033540	Homo sapiens mitofusin 1 (MFN1), transcript variant 1, mRNA
NM_005612	Homo sapiens RE1-silencing transcription factor (REST), mRNA
NM_007085	Homo sapiens follistatin-like 1 (FSTL1), mRNA
NM_000993	Homo sapiens ribosomal protein L31 (RPL31), mRNA
NM_012180	Homo sapiens F-box only protein 8 (FBXO8), mRNA
NM_033182	Homo sapiens F-box protein FBX30 (FBX30), mRNA
NM_033406	Homo sapiens F-box only protein 3 (FBXO3), transcript variant 2, mRNA
NM_012175	Homo sapiens F-box only protein 3 (FBXO3), transcript variant 1, mRNA
NM_017425	Homo sapiens sperm autoantigenic protein 17 (SPA17), mRNA
NM_005633	Homo sapiens son of sevenless homolog 1 (Drosophila) (SOS1), mRNA
NM_003333	Homo sapiens ubiquitin A-52 residue ribosomal protein fusion product 1 (UBA52), mRNA
NM_019894	Homo sapiens transmembrane protease, serine 4 (TMPRSS4), mRNA
NM_033313	Homo sapiens CDC14 cell division cycle 14 homolog A (S. cerevisiae)
	(CDC14A), transcript variant 3, mRNA
NM_033312	Homo sapiens CDC14 cell division cycle 14 homolog A (S. cerevisiae)
ND4 000 (72	(CDC14A), transcript variant 2, mRNA
NM_003672	Homo sapiens CDC14 cell division cycle 14 homolog A (S. cerevisiae)

	(CDC14A), transcript variant 1, mRNA
ND4 005796	Homo sapiens serologically defined colon cancer antigen 33 (SDCCAG33),
NM_005786	mRNA
ND ( 002619	Homo sapiens mitogen-activated protein kinase kinase kinase kinase 3
NM_003618	, ,
ND 4 00 (577	(MAP4K3), mRNA
NM_006577	Homo sapiens UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase
\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\	1 (B3GNT1), transcript variant 1, mRNA
NM_020981	Homo sapiens UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide
	1 (B3GALT1), mRNA
NM_033252	Homo sapiens UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase
27.5 000054	1 (B3GNT1), transcript variant 2, mRNA
NM_002954	Homo sapiens ribosomal protein S27a (RPS27A), mRNA
NM_000971	Homo sapiens ribosomal protein L7 (RPL7), mRNA
NM_033344	Homo sapiens egl nine homolog 3 (C. elegans) (EGLN3), mRNA
NM_024023	Homo sapiens unkempt-like (Drosophila) (UNKL), mRNA
NM_033221	Homo sapiens tripartite motif-containing 14 (TRIM14), transcript variant 4,
	mRNA
NM_033220	Homo sapiens tripartite motif-containing 14 (TRIM14), transcript variant 3,
	mRNA
NM_033219	Homo sapiens tripartite motif-containing 14 (TRIM14), transcript variant 2,
	mRNA
NM_014788	Homo sapiens tripartite motif-containing 14 (TRIM14), transcript variant 1,
	mRNA
NM_006074	Homo sapiens tripartite motif-containing 22 (TRIM22), mRNA
NM_012210	Homo sapiens tripartite motif-containing 32 (TRIM32), mRNA
NM_007276	Homo sapiens chromobox homolog 3 (HP1 gamma homolog, Drosophila)
	(CBX3), mRNA
NM_025227	Homo sapiens hypothetical protein dJ726C3.2 (DJ726C3.2), mRNA
NM_015271	Homo sapiens tripartite motif-containing 2 (TRIM2), mRNA
NM_017838	Homo sapiens nucleolar protein family A, member 2 (H/ACA small nucleolar RNPs) (NOLA2), mRNA
NM_032993	Homo sapiens nucleolar protein family A, member 1 (H/ACA small nucleolar
	RNPs) (NOLA1), transcript variant 2, mRNA
NM_018983	Homo sapiens nucleolar protein family A, member 1 (H/ACA small nucleolar
	RNPs) (NOLA1), transcript variant 1, mRNA
NM_004722	Homo sapiens adaptor-related protein complex 4, mu 1 subunit (AP4M1),
	mRNA
NM_033066	Homo sapiens membrane protein, palmitoylated 4 (MAGUK p55 subfamily
	member 4) (MPP4), mRNA
NM_033030	Homo sapiens bol, boule-like (Drosophila) (BOLL), mRNA
NM_004216	Homo sapiens death effector domain-containing (DEDD), transcript variant 2, mRNA
NM 032998	Homo sapiens death effector domain-containing (DEDD), transcript variant 1,
	mRNA
NM 033010	Homo sapiens poly(rC) binding protein 4 (PCBP4), transcript variant 4, mRNA
NM 033009	Homo sapiens poly(rC) binding protein 4 (PCBP4), transcript variant 2, mRNA
NM 033008	Homo sapiens poly(rC) binding protein 4 (PCBP4), transcript variant 3, mRNA
NM 020418	Homo sapiens poly(rC) binding protein 4 (PCBP4), transcript variant 1, mRNA
NM 032944	Homo sapiens serine/threonine kinase 31 (STK31), transcript variant 2, mRNA
NM 031414	Homo sapiens serine/threonine kinase 31 (STK31), transcript variant 1, mRNA
NM 014302	Homo sapiens Sec61 gamma (SEC61G), mRNA
NM 013336	Homo sapiens protein transport protein SEC61 alpha subunit isoform 1
	The state of the s

	(CECC(1A1) PALA
NM 031431	(SEC61A1), mRNA
NM_015490	Homo sapiens tethering factor SEC34 (SEC34), mRNA
NM 004892	Homo sapiens secretory pathway component Sec31B-1 (SEC31B-1), mRNA
	Homo sapiens SEC22 vesicle trafficking protein-like 1 (S. cerevisiae) (SEC22L1), mRNA
NM_032970	Homo sapiens vesicle trafficking protein (SEC22C), transcript variant 1, mRNA
NM_000969	Homo sapiens ribosomal protein L5 (RPL5), mRNA
NM_005034	Homo sapiens polymerase (RNA) II (DNA directed) polypeptide K (7.0kD) (POLR2K), mRNA
NM 014459	Homo sapiens protocadherin 17 (PCDH17), mRNA
NM_032961	Homo sapiens protocadherin 10 (PCDH10), transcript variant 1, mRNA
NM_020815	Homo sapiens protocadherin 10 (PCDH10), transcript variant 2, mRNA
NM_031988	Homo sapiens mitogen-activated protein kinase kinase 6 (MAP2K6), transcript variant 2, mRNA
NM_002758	Homo sapiens mitogen-activated protein kinase kinase 6 (MAP2K6), transcript variant 1, mRNA
NM_032419	Homo sapiens dom-3 homolog Z (C. elegans) (DOM3Z), transcript variant 1, mRNA
NM 032966	
	Homo sapiens Burkitt lymphoma receptor 1, GTP binding protein (BLR1), transcript variant 2, mRNA
NM_001716	Homo sapiens Burkitt lymphoma receptor 1, GTP binding protein (BLR1), transcript variant 1, mRNA
NM_004951	Homo sapiens Epstein-Barr virus induced gene 2 (lymphocyte-specific G
	protein-coupled receptor) (EBI2), mRNA
NM_004874	Homo sapiens BCL2-associated athanogene 4 (BAG4), mRNA
NM_001016	Homo sapiens ribosomal protein S12 (RPS12), mRNA
NM_031994	Homo sapiens ring finger protein 17 (RNF17), transcript variant short, mRNA
NM_031271	Homo sapiens testis expressed sequence 15 (TEX15), mRNA
NM_018995	Homo sapiens Mov1011, Moloney leukemia virus 10-like 1, homolog (mouse) (MOV10L1), mRNA
NM_032510	Homo sapiens par-6 partitioning defective 6 homolog gamma (C. elegans) (PARD6G), mRNA
NM_006704	Homo sapiens suppressor of G2 allele of SKP1, S. cerevisiae, homolog of (SGT1), mRNA
NM_031968	Homo sapiens nuclear prelamin A recognition factor (NARF), transcript variant 2, mRNA
NM_012336	Homo sapiens nuclear prelamin A recognition factor (NARF), transcript variant 1, mRNA
NM_003980	Homo sapiens microtubule-associated protein 7 (MAP7), mRNA
NM_032380	Homo sapiens elongation factor G2 (EFG2), mRNA
NM_032214	Homo sapiens Src-like-adaptor 2 (SLA2), mRNA
NM_020064	Homo sapiens BarH-like 1 (Drosophila) (BARHL1), mRNA
NM_005916	Homo sapiens MCM7 minichromosome maintenance deficient 7 (S. cerevisiae)
NM_004098	(MCM7), mRNA
NM_005826	Homo sapiens empty spiracles homolog 2 (Drosophila) (EMX2), mRNA
NM_006418	Homo sapiens differentially and the Homo sapiens differentially an
	Homo sapiens differentially expressed in hematopoietic lineages (GW112), mRNA
NM_005016	Homo sapiens poly(rC) binding protein 2 (PCBP2), transcript variant 1, mRNA
NM_031989	Homo sapiens poly(rC) binding protein 2 (PCBP2), transcript variant 2, mRNA
NM_006196	Homo sapiens poly(rC) binding protein 1 (PCBP1), mRNA
NM_031844	Homo sapiens heterogeneous nuclear ribonucleoprotein U (scaffold attachment

	factor A) (HNRPU), transcript variant 1, mRNA
NM_004501	Homo sapiens heterogeneous nuclear ribonucleoprotein U (scaffold attachment
	factor A) (HNRPU), transcript variant 2, mRNA
NM_004500	Homo sapiens heterogeneous nuclear ribonucleoprotein C (C1/C2) (HNRPC),
	transcript variant 2, mRNA
NM_031314	Homo sapiens heterogeneous nuclear ribonucleoprotein C (C1/C2) (HNRPC),
	transcript variant 1, mRNA
NM_031370	Homo sapiens heterogeneous nuclear ribonucleoprotein D (AU-rich element
	RNA binding protein 1, 37kD) (HNRPD), transcript variant 1, mRNA
NM_031369	Homo sapiens heterogeneous nuclear ribonucleoprotein D (AU-rich element
	RNA binding protein 1, 37kD) (HNRPD), transcript variant 2, mRNA
NM_002138	Homo sapiens heterogeneous nuclear ribonucleoprotein D (AU-rich element
	RNA binding protein 1, 37kD) (HNRPD), transcript variant 3, mRNA
NM_003903	Homo sapiens CDC16 cell division cycle 16 homolog (S. cerevisiae) (CDC16),
_	mRNA
NM_031483	Homo sapiens itchy homolog E3 ubiquitin protein ligase (mouse) (ITCH),
_	mRNA
NM 031907	Homo sapiens ubiquitin specific protease 26 (USP26), mRNA
NM 031866	Homo sapiens frizzled homolog 8 (Drosophila) (FZD8), mRNA
NG 000004	Homo sapiens genomic cytochrome P450, subfamily IIIA (niphedipine oxidase)
	(CYP3A) on chromosome 7
NM 001788	Homo sapiens CDC10 cell division cycle 10 homolog (S. cerevisiae) (CDC10),
	mRNA
NM 004276	Homo sapiens calcium binding protein 1 (calbrain) (CABP1), transcript variant
	2, mRNA
NM_031205	Homo sapiens calcium binding protein 1 (calbrain) (CABP1), transcript variant
	1, mRNA
NM_000784	Homo sapiens cytochrome P450, subfamily XXVIIA (steroid 27-hydroxylase,
_	cerebrotendinous xanthomatosis), polypeptide 1 (CYP27A1), nuclear gene
	encoding mitochondrial protein, mRNA
NM_031491	Homo sapiens retinol binding protein 5, cellular (RBP5), mRNA
NM_006929	Homo sapiens superkiller viralicidic activity 2-like (S. cerevisiae) (SKIV2L),
_	mRNA
NM 001447	Homo sapiens FAT tumor suppressor homolog 2 (Drosophila) (FAT2), mRNA
NM 007242	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 19 (DBP5
-	homolog, yeast) (DDX19), mRNA
NM_006773	
_	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 18 (Myc-
	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 18 (Myc-regulated) (DDX18), mRNA
NM 030655	regulated) (DDX18), mRNA
NM_030655	regulated) (DDX18), mRNA Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 11 (CHL1-like
	regulated) (DDX18), mRNA  Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 11 (CHL1-like helicase homolog, S. cerevisiae) (DDX11), transcript variant 3, mRNA
NM_030655 NM_030653	regulated) (DDX18), mRNA  Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 11 (CHL1-like helicase homolog, S. cerevisiae) (DDX11), transcript variant 3, mRNA  Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 11 (CHL1-like
NM_030653	regulated) (DDX18), mRNA  Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 11 (CHL1-like helicase homolog, S. cerevisiae) (DDX11), transcript variant 3, mRNA  Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 11 (CHL1-like helicase homolog, S. cerevisiae) (DDX11), transcript variant 1, mRNA
	regulated) (DDX18), mRNA  Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 11 (CHL1-like helicase homolog, S. cerevisiae) (DDX11), transcript variant 3, mRNA  Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 11 (CHL1-like helicase homolog, S. cerevisiae) (DDX11), transcript variant 1, mRNA  Homo sapiens cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase),
NM_030653 NM_000770	regulated) (DDX18), mRNA  Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 11 (CHL1-like helicase homolog, S. cerevisiae) (DDX11), transcript variant 3, mRNA  Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 11 (CHL1-like helicase homolog, S. cerevisiae) (DDX11), transcript variant 1, mRNA  Homo sapiens cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase), polypeptide 8 (CYP2C8), transcript variant Hp1-1, mRNA
NM_030653	regulated) (DDX18), mRNA  Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 11 (CHL1-like helicase homolog, S. cerevisiae) (DDX11), transcript variant 3, mRNA  Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 11 (CHL1-like helicase homolog, S. cerevisiae) (DDX11), transcript variant 1, mRNA  Homo sapiens cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase), polypeptide 8 (CYP2C8), transcript variant Hp1-1, mRNA  Homo sapiens cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase),
NM_030653 NM_000770 NM_030878	regulated) (DDX18), mRNA  Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 11 (CHL1-like helicase homolog, S. cerevisiae) (DDX11), transcript variant 3, mRNA  Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 11 (CHL1-like helicase homolog, S. cerevisiae) (DDX11), transcript variant 1, mRNA  Homo sapiens cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase), polypeptide 8 (CYP2C8), transcript variant Hp1-1, mRNA  Homo sapiens cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase), polypeptide 8 (CYP2C8), transcript variant Hp1-2, mRNA
NM_030653 NM_000770	regulated) (DDX18), mRNA  Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 11 (CHL1-like helicase homolog, S. cerevisiae) (DDX11), transcript variant 3, mRNA  Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 11 (CHL1-like helicase homolog, S. cerevisiae) (DDX11), transcript variant 1, mRNA  Homo sapiens cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase), polypeptide 8 (CYP2C8), transcript variant Hp1-1, mRNA  Homo sapiens cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase), polypeptide 8 (CYP2C8), transcript variant Hp1-2, mRNA  Homo sapiens sirtuin silent mating type information regulation 2 homolog 3 (S.
NM_030653 NM_000770 NM_030878 NM_012239	regulated) (DDX18), mRNA  Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 11 (CHL1-like helicase homolog, S. cerevisiae) (DDX11), transcript variant 3, mRNA  Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 11 (CHL1-like helicase homolog, S. cerevisiae) (DDX11), transcript variant 1, mRNA  Homo sapiens cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase), polypeptide 8 (CYP2C8), transcript variant Hp1-1, mRNA  Homo sapiens cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase), polypeptide 8 (CYP2C8), transcript variant Hp1-2, mRNA  Homo sapiens sirtuin silent mating type information regulation 2 homolog 3 (S. cerevisiae) (SIRT3), mRNA
NM_030653 NM_000770 NM_030878	regulated) (DDX18), mRNA  Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 11 (CHL1-like helicase homolog, S. cerevisiae) (DDX11), transcript variant 3, mRNA  Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 11 (CHL1-like helicase homolog, S. cerevisiae) (DDX11), transcript variant 1, mRNA  Homo sapiens cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase), polypeptide 8 (CYP2C8), transcript variant Hp1-1, mRNA  Homo sapiens cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase), polypeptide 8 (CYP2C8), transcript variant Hp1-2, mRNA  Homo sapiens sirtuin silent mating type information regulation 2 homolog 3 (S. cerevisiae) (SIRT3), mRNA  Homo sapiens sirtuin silent mating type information regulation 2 homolog 2 (S.
NM_030653 NM_000770 NM_030878 NM_012239	regulated) (DDX18), mRNA  Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 11 (CHL1-like helicase homolog, S. cerevisiae) (DDX11), transcript variant 3, mRNA  Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 11 (CHL1-like helicase homolog, S. cerevisiae) (DDX11), transcript variant 1, mRNA  Homo sapiens cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase), polypeptide 8 (CYP2C8), transcript variant Hp1-1, mRNA  Homo sapiens cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase), polypeptide 8 (CYP2C8), transcript variant Hp1-2, mRNA  Homo sapiens sirtuin silent mating type information regulation 2 homolog 3 (S. cerevisiae) (SIRT3), mRNA

NM_012238	Homo sapiens sirtuin silent mating type information regulation 2 homolog 1 (S. cerevisiae) (SIRT1), mRNA
NM_031309	Homo sapiens scratch homolog 1, zinc finger protein (Drosophila) (SCRT1),
37.5	mRNA
NM_031278	Homo sapiens tudor domain containing 1 (TDRD1), mRNA
NM_031277	Homo sapiens ring finger protein 17 (RNF17), transcript variant long, mRNA
NM_031276	Homo sapiens testis expressed sequence 11 (TEX11), mRNA
NM_031273	Homo sapiens testis expressed sequence 13B (TEX13B), mRNA
NM_031272	Homo sapiens testis expressed sequence 14 (TEX14), mRNA
NM_006636	Homo sapiens methylene tetrahydrofolate dehydrogenase (NAD+ dependent), methenyltetrahydrofolate cyclohydrolase (MTHFD2), nuclear gene encoding
NM_022818	mitochondrial protein, mRNA  Homo sapiens microtubule-associated proteins 1A/1B light chain 3
	(MAP1A/1BLC3), mRNA
NM_018607	Homo sapiens hypothetical protein PRO1853 (PRO1853), mRNA
NM 004856	Homo sapiens kinesin-like 5 (mitotic kinesin-like protein 1) (KNSL5), mRNA
NM_030979	Homo sapiens poly(A) binding protein, cytoplasmic 3 (PABPC3), mRNA
NM 030770	Homo sapiens transmembrane protease, serine 5 (spinesin) (TMPRSS5), mRNA
NM 002545	Homo sapiens opioid binding protein/cell adhesion molecule-like (OPCML),
	mRNA
NM_014676	Homo sapiens pumilio homolog 1 (Drosophila) (PUM1), mRNA
NM_030673	Homo sapiens SEC13-like 1 (S. cerevisiae) (SEC13L1), mRNA
NM_003342	Homo sapiens ubiquitin-conjugating enzyme E2G 1 (UBC7 homolog, C.
77.5	elegans) (UBE2G1), mRNA
NM_022051	Homo sapiens egl nine homolog 1 (C. elegans) (EGLN1), mRNA
NM_015577	Homo sapiens retinoic acid induced 14 (RAI14), mRNA
NM_012170	Homo sapiens F-box only protein 22 (FBXO22), mRNA
NM_022304	Homo sapiens histamine receptor H2 (HRH2), mRNA
NM_022333	Homo sapiens TIA1 cytotoxic granule-associated RNA binding protein-like 1 (TIAL1), transcript variant 2, mRNA
NM_003252	Homo sapiens TIA1 cytotoxic granule-associated RNA binding protein-like 1
	(TIAL1), transcript variant 1, mRNA
NM_017910	Homo sapiens hypothetical protein FLJ20628 (FLJ20628), mRNA
NM_012384	Homo sapiens glucocorticoid modulatory element binding protein 2 (GMEB2), mRNA
NM_006118	Homo sapiens HS1 binding protein (HAX1), mRNA
NM_022740	Homo sapiens homeodomain interacting protein kinase 2 (HIPK2), mRNA
NM_002005	Homo sapiens feline sarcoma oncogene (FES), mRNA
NM_014757	Homo sapiens mastermind-like 1 (Drosophila) (MAML1), mRNA
NM_025136	Homo sapiens optic atrophy 3 (autosomal recessive, with chorea and spastic paraplegia) (OPA3), mRNA
NM_024505	Homo sapiens NADPH oxidase, EF hand calcium-binding domain 5 (NOX5), mRNA
NM_022362	Homo sapiens MMS19-like (MET18 homolog, S. cerevisiae) (MMS19L), mRNA
NM_000256	Homo sapiens myosin binding protein C, cardiac (MYBPC3), mRNA
NM_000276	Homo sapiens oculocerebrorenal syndrome of Lowe (OCRL), transcript variant a, mRNA
NM_001587	Homo sapiens oculocerebrorenal syndrome of Lowe (OCRL), transcript variant b, mRNA
NM_001407	Homo sapiens cadherin, EGF LAG seven-pass G-type receptor 3 (flamingo homolog, Drosophila) (CELSR3), mRNA
	, \/,

NM_001408	Homo sapiens cadherin, EGF LAG seven-pass G-type receptor 2 (flamingo
_	homolog, Drosophila) (CELSR2), mRNA
NM_005735	Homo sapiens ARP1 actin-related protein 1 homolog B, centractin beta (yeast) (ACTR1B), mRNA
NM_012254	Homo sapiens very long-chain acyl-CoA synthetase homolog 2 (VLCS-H2), mRNA
NM_012331	Homo sapiens methionine sulfoxide reductase A (MSRA), mRNA
NM_016596	Homo sapiens histone deacetylase 7A (HDAC7A), transcript variant 2, mRNA
NM_015401	Homo sapiens histone deacetylase 7A (HDAC7A), transcript variant 1, mRNA
NM_004082	Homo sapiens dynactin 1 (p150, glued homolog, Drosophila) (DCTN1), transcript variant 1, mRNA
NM_023019	Homo sapiens dynactin 1 (p150, glued homolog, Drosophila) (DCTN1), transcript variant 2, mRNA
NM 002893	Homo sapiens retinoblastoma binding protein 7 (RBBP7), mRNA
NM_023001	Homo sapiens retinoblastoma binding protein 1 (RBBP1), transcript variant 3, mRNA
NM_023000	Homo sapiens retinoblastoma binding protein 1 (RBBP1), transcript variant 2, mRNA
NM_002892	Homo sapiens retinoblastoma binding protein 1 (RBBP1), transcript variant 1, mRNA
NM_024408	Homo sapiens Notch homolog 2 (Drosophila) (NOTCH2), mRNA
NM_012311	Homo sapiens KIN, antigenic determinant of recA protein homolog (mouse) (KIN), mRNA
NM_021938	Homo sapiens bruno-like 5, RNA binding protein (Drosophila) (BRUNOL5), mRNA
NM_020180	Homo sapiens bruno-like 4, RNA binding protein (Drosophila) (BRUNOL4), mRNA
NM_005868	Homo sapiens BET1 homolog (S. cerevisiae) (BET1), mRNA
NM_002467	Homo sapiens v-myc myelocytomatosis viral oncogene homolog (avian) (MYC), mRNA
NM_022817	Homo sapiens period homolog 2 (Drosophila) (PER2), transcript variant 1, mRNA
NM_003894	Homo sapiens period homolog 2 (Drosophila) (PER2), transcript variant 2, mRNA
NM_006660	Homo sapiens ClpX caseinolytic protease X homolog (E. coli) (CLPX), mRNA
NM_012394	Homo sapiens prefoldin 2 (PFDN2), mRNA
NM_004234	Homo sapiens zinc finger protein 93 homolog (mouse) (ZFP93), mRNA
NM_005870	Homo sapiens sin3-associated polypeptide, 18kD (SAP18), mRNA
NM_003350	Homo sapiens ubiquitin-conjugating enzyme E2 variant 2 (UBE2V2), mRNA
NM_022476	Homo sapiens fused toes homolog (mouse) (FTS), mRNA
NM_022444	Homo sapiens solute carrier family 13 (sodium/sulfate symporters), member 1 (SLC13A1), mRNA
NM_018127	Homo sapiens elaC homolog 2 (E. coli) (ELAC2), mRNA
NM_014317	Homo sapiens trans-prenyltransferase (TPT), mRNA
NM_022173	Homo sapiens TIA1 cytotoxic granule-associated RNA binding protein (TIA1), transcript variant 2, mRNA
NM_022037	Homo sapiens TIA1 cytotoxic granule-associated RNA binding protein (TIA1), transcript variant 1, mRNA
NM_004973	Homo sapiens jumonji homolog (mouse) (JMJ), mRNA
NM_021971	Homo sapiens GDP-mannose pyrophosphorylase B (GMPPB), transcript variant 2, mRNA
NM_013334	Homo sapiens GDP-mannose pyrophosphorylase B (GMPPB), transcript variant

	1 DNIA
NM_013335	1, mRNA
	Homo sapiens GDP-mannose pyrophosphorylase A (GMPPA), mRNA
NM_021267	Homo sapiens LAG1 longevity assurance homolog 1 (S. cerevisiae) (LASS1), mRNA
NM_005811	Homo sapiens growth differentiation factor 11 (GDF11), mRNA
NM_005971	Homo sapiens FXYD domain-containing ion transport regulator 3 (FXYD3), transcript variant 1, mRNA
NM_021910	Homo sapiens FXYD domain-containing ion transport regulator 3 (FXYD3), transcript variant 2, mRNA
NM_022096	Homo sapiens ankyrin repeat domain 5 (ANKRD5), mRNA
NM_022073	Homo sapiens egl nine homolog 3 (C. elegans) (EGLN3), mRNA
NM_022047	Homo sapiens differentially expressed in FDCP 6 homolog (mouse) (DEF6), mRNA
NM_021778	Homo sapiens a disintegrin and metalloproteinase domain 28 (ADAM28), transcript variant 2, mRNA
NM_021777	Homo sapiens a disintegrin and metalloproteinase domain 28 (ADAM28), transcript variant 3, mRNA
NM_000152	Homo sapiens glucosidase, alpha; acid (Pompe disease, glycogen storage disease type II) (GAA), mRNA
NM_002910	Homo sapiens renin binding protein (RENBP), mRNA
NM_012072	Homo sapiens complement component 1, q subcomponent, receptor 1 (C1QR1), mRNA
NM_000534	Homo sapiens PMS1 postmeiotic segregation increased 1 (S. cerevisiae) (PMS1), mRNA
NM_005451	Homo sapiens enigma (LIM domain protein) (ENIGMA), mRNA
NM_021975	Homo sapiens v-rel reticuloendotheliosis viral oncogene homolog A, nuclear
	factor of kappa light polypeptide gene enhancer in B-cells 3, p65 (avian) (RELA), mRNA
NM_021958	Homo sapiens H2.0-like homeo box 1 (Drosophila) (HLX1), mRNA
NM_004139	Homo sapiens lipopolysaccharide binding protein (LBP), mRNA
NM_005442	Homo sapiens eomesodermin homolog (Xenopus laevis) (EOMES), mRNA
NM_004187	Homo sapiens Smcx homolog, X chromosome (mouse) (SMCX), mRNA
NM_003170	Homo sapiens suppressor of Ty 6 homolog (S. cerevisiae) (SUPT6H), mRNA
NM_003062	Homo sapiens slit homolog 3 (Drosophila) (SLIT3), mRNA
NM_003068	Homo sapiens slug homolog, zinc finger protein (chicken) (SLUG), mRNA
NM_021824	Homo sapiens NIF3 NGG1 interacting factor 3-like 1 (S. pombe) (NIF3L1), mRNA
NM_021783	Homo sapiens ectodysplasin A2 isoform receptor (XEDAR), mRNA
NM_004196	Homo sapiens cyclin-dependent kinase-like 1 (CDC2-related kinase) (CDKL1), mRNA
NM_000535	Homo sapiens PMS2 postmeiotic segregation increased 2 (S. cerevisiae) (PMS2), mRNA
NM_002356	Homo sapiens myristoylated alanine-rich protein kinase C substrate (MARCKS), mRNA
NM_021728	Homo sapiens orthodenticle homolog 2 (Drosophila) (OTX2), mRNA
NM_014588	Homo sapiens visual system homeobox 1 homolog, CHX10-like (zebrafish) (VSX1), mRNA
NM_003503	Homo sapiens CDC7 cell division cycle 7-like 1 (S. cerevisiae) (CDC7L1), mRNA
NM_004059	Homo sapiens cysteine conjugate-beta lyase; cytoplasmic (glutamine transaminase K, kyneurenine aminotransferase) (CCBL1), mRNA
NM 020651	Homo sapiens pellino homolog 1 (Drosophila) (PELI1), mRNA
	The permit homolog i (Diosophila) (i DDII), IIIKNA

NM_018411	Homo sapiens hairless homolog (mouse) (HR), mRNA
NM_014569	Homo sapiens zinc finger protein 95 homolog (mouse) (ZFP95), mRNA
NM_012458	Homo sapiens translocase of inner mitochondrial membrane 13 homolog B
	(yeast) (TIMM13B), mRNA
NM_000672	Homo sapiens alcohol dehydrogenase 6 (class V) (ADH6), mRNA
NM_003603	Homo sapiens Arg/Abl-interacting protein ArgBP2 (ARGBP2), transcript variant
	1, mRNA
NM_021069	Homo sapiens Arg/Abl-interacting protein ArgBP2 (ARGBP2), transcript variant
	2, mRNA
NM_004950	Homo sapiens dermatan sulfate proteoglycan 3 (DSPG3), mRNA
NM_004701	Homo sapiens cyclin B2 (CCNB2), mRNA
NM_021100	Homo sapiens NFS1 nitrogen fixation 1 (S. cerevisiae) (NFS1), mRNA
NM_021255	Homo sapiens pellino homolog 2 (Drosophila) (PELI2), mRNA
NM_021115	Homo sapiens seizure related 6 homolog (mouse)-like (SEZ6L), mRNA
NM_004756	Homo sapiens numb homolog (Drosophila)-like (NUMBL), mRNA
NM_004690	Homo sapiens LATS, large tumor suppressor, homolog 1 (Drosophila) (LATS1), mRNA
NM_000461	Homo sapiens thyroid hormone receptor, beta (erythroblastic leukemia viral (verb-a) oncogene homolog 2, avian) (THRB), mRNA
NM_021078	Homo sapiens GCN5 general control of amino-acid synthesis 5-like 2 (yeast) (GCN5L2), mRNA
NM 002877	Homo sapiens RAD51-like 1 (S. cerevisiae) (RAD51L1), mRNA
NM_001552	Homo sapiens insulin-like growth factor binding protein 4 (IGFBP4), mRNA
NM 002487	Homo sapiens necdin homolog (mouse) (NDN), mRNA
NM 012425	Homo sapiens Ras suppressor protein 1 (RSU1), mRNA
NM_005618	Homo sapiens delta-like 1 (Drosophila) (DLL1), mRNA
NM_021038	Homo sapiens muscleblind-like (Drosophila) (MBNL), mRNA
NM_014268	Homo sapiens microtubule-associated protein, RP/EB family, member 2 (MAPRE2), mRNA
NM_020662	Homo sapiens MRS2-like, magnesium homeostasis factor (S. cerevisiae) (MRS2L), mRNA
NM_020649	Homo sapiens chromobox homolog 8 (Pc class homolog, Drosophila) (CBX8), mRNA
NM 018436	Homo sapiens allantoicase (ALLC), mRNA
NM 020528	Homo sapiens poly(rC) binding protein 3 (PCBP3), mRNA
NM_014276	Homo sapiens recombining binding protein suppressor of hairless (Drosophila)-like (RBPSUHL), mRNA
NM 019557	Homo sapiens hypothetical protein RP1-317E23 (LOC56181), mRNA
NM 020347	Homo sapiens leucine zipper transcription factor-like 1 (LZTFL1), mRNA
NM_005744	Homo sapiens ariadne homolog, ubiquitin-conjugating enzyme E2 binding protein, 1 (Drosophila) (ARIH1), mRNA
NM 007044	Homo sapiens katanin p60 (ATPase-containing) subunit A 1 (KATNA1), mRNA
NM 002688	Homo sapiens peanut-like 1 (Drosophila) (PNUTL1), mRNA
NM_013384	Homo sapiens LAG1 longevity assurance homolog 2 (S. cerevisiae) (LASS2), mRNA
NM 020230	Homo sapiens peter pan homolog (Drosophila) (PPAN), mRNA
NM_020182	Homo sapiens transmembrane, prostate androgen induced RNA (TMEPAI), mRNA
NM 020248	Homo sapiens catenin, beta interacting protein 1 (CTNNBIP1), mRNA
NM_000399	Homo sapiens early growth response 2 (Krox-20 homolog, Drosophila) (EGR2), mRNA
NM_002965	Homo sapiens S100 calcium binding protein A9 (calgranulin B) (S100A9),
·	· · · · · · · · · · · · · · · · · · ·

ND 6 000064	mRNA
NM_002964	Homo sapiens S100 calcium binding protein A8 (calgranulin A) (S100A8),
377.5.0000.60	MRNA
NM_002963	Homo sapiens S100 calcium binding protein A7 (psoriasin 1) (S100A7), mRNA
NM_014624	Homo sapiens \$100 calcium binding protein A6 (calcyclin) (\$100A6) mRNA
NM_019554	Homo sapiens \$100 calcium binding protein A4 (calcium protein, calvasculin
27.6 000061	metastasin, murine placental homolog) (S100A4), transcript variant 2 mRNA
NM_002961	Homo sapiens \$100 calcium binding protein A4 (calcium protein, calvasculin
) The 007070	metastasin, murine placental homolog) (S100A4), transcript variant 1 mRNA
NM_005978	Homo sapiens \$100 calcium binding protein A2 (\$100A2), mRNA
NM_002537	Homo sapiens ornithine decarboxylase antizyme 2 (OAZ2) mRNA
NM_019854	Homo sapiens HMT1 hnRNP methyltransferase-like 3 (S. cerevisiae)
ND 6 010610	(HRMT1L3), mRNA
NM_019619	Homo sapiens par-3 partitioning defective 3 homolog (C. elegans) (PARD3), mRNA
NM_017454	Homo sapiens staufen, RNA binding protein (Drosophila) (STAU), transcript
	variant 11, mr.NA
NM_017453	Homo sapiens staufen, RNA binding protein (Drosophila) (STAU), transcript
	Variant 13, mkNA
NM_017452	Homo sapiens staufen, RNA binding protein (Drosophila) (STAU), transcript
	Variant 12, mrnA
NM_003785	Homo sapiens G antigen, family B, 1 (prostate associated) (GAGEB1), mRNA
NM_015044	Homo sapiens golgi associated, gamma adaptin ear containing. ARF binding
	protein 2 (GGA2), mRNA
NM_013365	Homo sapiens golgi associated, gamma adaptin ear containing, ARF binding
	protein 1 (GGA1), mRNA
NM_004781	Homo sapiens vesicle-associated membrane protein 3 (cellubrevin) (VAMP3),
3.77.6.04.0.60.5	mrna
NM_018685	Homo sapiens anillin, actin binding protein (scraps homolog, Drosophila)
ND 6 017007	(ANLN), mRNA
NM_017927	Homo sapiens mitofusin 1 (MFN1), transcript variant 2, mRNA
NM 018387	Homo sapiens spermatid perinuclear RNA binding protein (STRRP) mRNA
NM_018378	Homo sapiens F-box and leucine-rich repeat protein 8 (FBXL8) mRNA
NM_018158	Homo sapiens solute carrier family 4 (anion exchanger), member 1, adaptor
NM 019022	protein (SLC4A1AP), mRNA
NM 018032	Homo sapiens LUC7-like (S. cerevisiae) (LUC7L), mRNA
NM_017575	Homo sapiens chromosome 17 open reading frame 31 (C17orf31), mRNA
NM_018696	Homo sapiens elaC homolog 1 (E. coli) (ELAC1), mRNA
NM_005781	Homo sapiens activated p21cdc42Hs kinase (ACK1), mRNA
NM_016831	Homo sapiens period homolog 3 (Drosophila) (PER3), mRNA
NM_003387	Homo sapiens Wiskott-Aldrich syndrome protein interacting protein (WASPIP), mRNA
NM_005993	Homo sapiens tubulin-specific chaperone d (TBCD), mRNA
NM_003014	Homo sapiens secreted frizzled-related protein 4 (SFRP4), mRNA
NM_006744	Homo sapiens retinol binding protein 4, plasma (RBP4), mRNA
NM_002899	Homo sapiens retinol binding protein 1, cellular (RBP1), mRNA
NM_005524	Homo sapiens hairy homolog (Drosophila) (HRY), mRNA
NM_005206	Homo sapiens v-crk sarcoma virus CT10 oncogene homolog (avian) (CRK),
	transcript variant I, mRNA
NM_016823	Homo sapiens v-crk sarcoma virus CT10 oncogene homolog (avian) (CRK),
	transcript variant II, mRNA
NM_016948	Homo sapiens par-6 partitioning defective 6 homolog alpha (C.elegans)
	b with (C.Cicgails)

	(PARD6A), mRNA
NM 017420	Homo sapiens sine oculis homeobox homolog 4 (Drosophila) (SIX4), mRNA
NM 016932	Homo sapiens sine oculis homeobox homolog 2 (Drosophila) (SIX2), mRNA
NM 017415	Homo sapiens kelch-like 3 (Drosophila) (KLHL3), mRNA
NM 017412	Homo sapiens frizzled homolog 3 (Drosophila) (FZD3), mRNA
NM 003400	Homo sapiens exportin 1 (CRM1 homolog, yeast) (XPO1), mRNA
NM 002889	Homo sapiens retinoic acid receptor responder (tazarotene induced) 2
1111_00200	(RARRES2), mRNA
NM_006064	Homo sapiens GTP-binding protein ragB (RAGB), transcript variant RAGBs, mRNA
NM_016656	Homo sapiens GTP-binding protein ragB (RAGB), transcript variant RAGBl, mRNA
NM_003857	Homo sapiens galanin receptor 2 (GALR2), mRNA
NM_016655	Homo sapiens GA binding protein transcription factor, beta subunit 2 (47kD) (GABPB2), transcript variant gamma, mRNA
NM_002041	Homo sapiens GA binding protein transcription factor, beta subunit 2 (47kD) (GABPB2), transcript variant gamma, mRNA
NM_016654	Homo sapiens GA binding protein transcription factor, beta subunit 1 (53kD) (GABPB1), transcript variant beta, mRNA
NM_005254	Homo sapiens GA binding protein transcription factor, beta subunit 1 (53kD) (GABPB1), transcript variant beta, mRNA
NM_015843	Homo sapiens LIM domain only 7 (LMO7), transcript variant 3, mRNA
NM_015842	Homo sapiens LIM domain only 7 (LMO7), transcript variant 2, mRNA
NM_002228	Homo sapiens v-jun sarcoma virus 17 oncogene homolog (avian) (JUN), mRNA
NM 016178	Homo sapiens ornithine decarboxylase antizyme 3 (OAZ3), mRNA
NM_016538	Homo sapiens sirtuin silent mating type information regulation 2 homolog 7 (S. cerevisiae) (SIRT7), mRNA
NM_016539	Homo sapiens sirtuin silent mating type information regulation 2 homolog 6 (S. cerevisiae) (SIRT6), mRNA
NM 016316	Homo sapiens REV1-like (yeast) (REV1L), mRNA
NM_016138	Homo sapiens COQ7 coenzyme Q, 7 homolog ubiquinone (yeast) (COQ7), mRNA
NM_016583	Homo sapiens palate, lung and nasal epithelium carcinoma associated (PLUNC), mRNA
NM_015886	Homo sapiens protease inhibitor 15 (PI15), mRNA
NM_016067	Homo sapiens mitochondrial ribosomal protein S18C (MRPS18C), nuclear gene encoding mitochondrial protein, mRNA
NM_015946	Homo sapiens pelota homolog (Drosophila) (PELO), mRNA
NM_016397	Homo sapiens TH1-like (Drosophila) (TH1L), mRNA
NM_016587	Homo sapiens chromobox homolog 3 (HP1 gamma homolog, Drosophila) (CBX3), mRNA
NM 016347	Homo sapiens putative N-acetyltransferase Camello 2 (CML2), mRNA
NM 015727	Homo sapiens tachykinin receptor 1 (TACR1), transcript variant short, mRNA
NM 001058	Homo sapiens tachykinin receptor 1 (TACR1), transcript variant long, mRNA
NM_004052	Homo sapiens BCL2/adenovirus E1B 19kD interacting protein 3 (BNIP3),
NM_014820	nuclear gene encoding mitochondrial protein, mRNA  Homo sapiens translocase of outer mitochondrial membrane 70 homolog A (yeast) (TOMM70A), mRNA
NM 014918	Homo sapiens carbohydrate (chondroitin) synthase 1 (CHSY1), mRNA
NM_014707	Homo sapiens histone deacetylase 9 (HDAC9-PENDING), transcript variant 3, mRNA
NM 014683	Homo sapiens unc-51-like kinase 2 (C. elegans) (ULK2), mRNA
	· · · · · · · · · · · · · · · · · · ·

NDA 014074	
NM_014874	Homo sapiens mitofusin 2 (MFN2), mRNA
NM 014071	Homo sapiens nuclear receptor coactivator 6 (NCOA6), mRNA
NM_015700	110110 Sapiells MIKA Interacting protein 5 (HIDIDS) -DATA
NM_015685	1101110 Sapiens Syndecan hinding protein (comtonin) 2 (CD CDD2)
NM_014263	Tiono sapiciis Tiviet-like I (S. cerevisiae) (VMF11 1) mDNIA
NM_014297	110110 Sapicils protein expressed in thuroid (VE121112) DATA
NM_014393	Homo sapiens staufen, RNA binding protein, homolog 2 (Drosophila) (STAU2)
NM_014403	Homo sapiens sialyltransferase 7D ((alpha-N-acetylneuraminyl-2,3-beta-
	galactosyl-1.3)-N-acetyl galactocominide -1.1. 2.6
	galactosyl-1,3)-N-acetyl galactosaminide alpha-2,6-sialyltransferase) (SIAT7D)
NM_014465	Homo sapiens sulfotransferase family, cytosolic, 1B, member 1 (SULT1B1),
	mRNA mRNA
NM 014485	
NM_014303	Homo sapiens prostaglandin D2 synthase, hematopoietic (PGDS), mRNA
	Homo sapiens pescadillo homolog 1, containing BRCT domain (zebrafish)  (PES1), mRNA
NM 014253	Homo saniens odz. odd Ozlawa 1 1 1 2
NM_014429	Homo sapiens microrabidio handle (ODZ1), mRNA
NM 006439	Tromo sapiens inicionalidia nomolog (monse) (MODC) -DNIA
NM_015322	Tiomo sapiens mao-21-like 2 (C. elegans) (MAR2112) mDNA
NM_014591	Tiomo sapiens iem-i nomolog h (C. elegans) (FEM1D) mDNIA
NM 004449	notice sapiens Ky channel interacting protein 2 (VCNID2) PALA
11111_004449	From Sapiens v-ets erythroblastosis virus F26 oncogene like (avior) (FDC)
ND4 014420	111111111
NM_014420	Homo sapiens dicklor fly and a Windowski (DKK4), mRNA
NM_014421	Tromo sapiens dickkopi nomojog / (Xenonis laevis) (DVV2) DVA
NM_014325	110110 Sapiciis Coronin, actin hinding protein 10 (CORO10) Park
NM_014246	Tiomo sapiens caunenn. E(if I A(i seven-nass G tyme recents 1/9
<u> </u>	1 Minorog, Diosophila) (CELSRI), mRNA
NM_014391	Homo sapiens cardiac ankyrin repeat protein (CARR) RALA
NM_014336	nomo sapiens aryl hydrocarbon receptor interacting protein like 1 (A DL 1)
NM_014265	Homo sapiens a disintegrin and metalloproteinase domain 28 (ADAM28),
	tanscript variant 1. Inkna
NM_014237	Homo sapiens a disintegrin and metalloproteinase domain 18 (ADAM18),
	mRNA (ADAM18),
NM_005032	Homo sapiens plastin 3 (T isoform) (PLS3), mRNA
NM_013980	Homo sapiens BCL2/adenovirus E1B 19kD interacting protein 1 (BNIP1),
	transcript variant BNIP1-c, mRNA
NM_013979	Homo sapiens BCL2/adenovirus E1B 19kD interacting protein 1 (BNIP1),
	transcript variant BNIP1-b, mRNA
NM_013978	Homo sapiens BCL2/adenovirus E1B 19kD interacting protein 1 (BNIP1),
	transcript variant BNIP1-a, mRNA
NM_004178	Homo sapiens TAR (HIV) DNA hinding
NM_005915	Homo sapiens TAR (HIV) RNA binding protein 2 (TARBP2), mRNA
	Homo sapiens MCM6 minichromosome maintenance deficient 6 (MIS5 homolog S, nombe) (S, acrossinics) (MCM6) (100 MIS5)
NM_002576	homolog, S. pombe) (S. cerevisiae) (MCM6), mRNA
	Homo sapiens p21/Cdc42/Rac1-activated kinase 1 (STE20 homolog, yeast) (PAK1), mRNA
NM_012091	(17AKI), IIIKWA
	Homo sapiens adenosine deaminase, tRNA-specific 1 (ADAT1), mRNA
005550	fromo sapiens Livi domain only 7 (LMO7) mRNA
	Homo sapiens fer-1-like 3, myoferlin (C. elegans) (FFR II 3) mpnia
	II-
NM_006113	Homo sapiens vav 3 oncogene (VAV3), mRNA  Homo sapiens carboxylesterase 2 (intestine, liver) (CES2), mRNA

NM_005721	Homo sapiens ARP3 actin-related protein 3 homolog (yeast) (ACTR3), mRNA
NM_003325	Homo sapiens HIR histone cell cycle regulation defective homolog A (S.
	cerevisiae) (HIRA), mRNA
NM_012242	Homo sapiens dickkopf homolog 1 (Xenopus laevis) (DKK1), mRNA
NM_012429	Homo sapiens SEC14-like 2 (S. cerevisiae) (SEC14L2), mRNA
NM 012190	Homo sapiens formyltetrahydrofolate dehydrogenase (FTHFD), mRNA
NM 005069	Homo sapiens single-minded homolog 2 (Drosophila) (SIM2), transcript variant
	SIM2, mRNA
NM 009586	Homo sapiens single-minded homolog 2 (Drosophila) (SIM2), transcript variant
	SIM2s, mRNA
NM 002610	Homo sapiens pyruvate dehydrogenase kinase, isoenzyme 1 (PDK1), nuclear
	gene encoding mitochondrial protein, mRNA
NM 013374	Homo sapiens programmed cell death 6 interacting protein (PDCD6IP), mRNA
NM 013367	Homo sapiens anaphase-promoting complex subunit 4 (APC4), mRNA
NM 002968	Homo sapiens sal-like 1 (Drosophila) (SALL1), mRNA
<del></del>	
NM_002449	Homo sapiens msh homeo box homolog 2 (Drosophila) (MSX2), mRNA
NM_006739	Homo sapiens MCM5 minichromosome maintenance deficient 5, cell division cycle 46 (S. cerevisiae) (MCM5), mRNA
NM_012460	Homo sapiens translocase of inner mitochondrial membrane 9 homolog (yeast)
_	(TIMM9), mRNA
NM_012457	Homo sapiens translocase of inner mitochondrial membrane 13 homolog A
_	(yeast) (TIMM13A), mRNA
NM_012456	Homo sapiens translocase of inner mitochondrial membrane 10 homolog (yeast)
_	(TIMM10), mRNA
NM 012450	Homo sapiens solute carrier family 13 (sodium/sulfate symporters), member 4
	(SLC13A4), mRNA
NM_012444	Homo sapiens SPO11 meiotic protein covalently bound to DSB-like (S.
	cerevisiae) (SPO11), mRNA
NM_012240	Homo sapiens sirtuin silent mating type information regulation 2 homolog 4 (S.
1111_012210	cerevisiae) (SIRT4), mRNA
NM 012387	Homo sapiens peptidyl arginine deiminase, type V (PAD), mRNA
NM 012381	Homo sapiens origin recognition complex, subunit 3-like (yeast) (ORC3L),
1414_012561	mRNA
NM_012225	Homo sapiens nucleotide binding protein 2 (MinD homolog, E. coli) (NUBP2),
	mRNA
NM_012222	Homo sapiens mutY homolog (E. coli) (MUTYH), mRNA
NM 012279	Homo sapiens double-stranded RNA-binding zinc finger protein JAZ (JAZ),
	mRNA
NM 012206	Homo sapiens hepatitis A virus cellular receptor 1 (HAVCR-1), mRNA
NM 012205	Homo sapiens 3-hydroxyanthranilate 3,4-dioxygenase (HAAO), mRNA
NM 012198	Homo sapiens grancalcin, EF-hand calcium binding protein (GCA), mRNA
NM 012193	Homo sapiens frizzled homolog 4 (Drosophila) (FZD4), mRNA
NM 012192	Homo sapiens fracture callus 1 homolog (rat) (FXC1), mRNA
NM_012076	Homo sapiens crumbs homolog 1 (Drosophila) (CRB1), mRNA
NM 012124	Homo sapiens cysteine and histidine-rich domain (CHORD)-containing, zinc
_	binding protein 1 (CHORDC1), mRNA
NM_012118	Homo sapiens CCR4 carbon catabolite repression 4-like (S. cerevisiae)
	(CCRN4L), mRNA
NM_012117	Homo sapiens chromobox homolog 5 (HP1 alpha homolog, Drosophila) (CBX5),
	mRNA
L	mid (1)
NM_012108	Homo sapiens BCR downstream signaling 1 (BRDG1), mRNA

NM_012094	Homo sapiens peroxiredoxin 5 (PRDX5), mRNA
NM_004506	Homo sapiens heat shock transcription factor 2 (HSF2), mRNA
NM 004423	Homo sapiens dishevelled, dsh homolog 3 (Drosophila) (DVL3), mRNA
NM 007374	Homo sapiens sine oculis homeobox homolog 6 (Drosophila (SIX6), mRNA
NM_007373	Homo sapiens soc-2 suppressor of clear homolog (C. elegans) (SHOC2), mRNA
NM_002388	Homo sapiens MCM3 minichromosome maintenance deficient 3 (S. cerevisiae)
_	(MCM3), mRNA
NM_004873	Homo sapiens BCL2-associated athanogene 5 (BAG5), mRNA
NM_007316	Homo sapiens agouti related protein homolog (mouse) (AGRP), transcript variant 2, mRNA
NM_003819	Homo sapiens poly(A) binding protein, cytoplasmic 4 (inducible form) (PABPC4), mRNA
NM_005737	Homo sapiens ADP-ribosylation factor-like 7 (ARL7), mRNA
NM_002358	Homo sapiens MAD2 mitotic arrest deficient-like 1 (yeast) (MAD2L1), mRNA
NM 007264	Homo sapiens adrenomedullin receptor (ADMR), mRNA
NM_006870	Homo sapiens destrin (actin depolymerizing factor) (DSTN), mRNA
NM_005476	Homo sapiens UDP-N-acetylglucosamine-2-epimerase/N-acetylmannosamine
	kinase (GNE), mRNA
NM_007309	Homo sapiens diaphanous homolog 2 (Drosophila) (DIAPH2), transcript variant 12C, mRNA
NM_001878	Homo sapiens cellular retinoic acid binding protein 2 (CRABP2), mRNA
NM_000489	Homo sapiens alpha thalassemia/mental retardation syndrome X-linked (RAD54
	homolog, S. cerevisiae) (ATRX), mRNA
NM_002528	Homo sapiens nth endonuclease III-like 1 (E. coli) (NTHL1), mRNA
NM_004085	Homo sapiens translocase of inner mitochondrial membrane 8 homolog A (yeast)
	(TIMM8A), nuclear gene encoding mitochondrial protein, mRNA
NM_002310	Homo sapiens leukemia inhibitory factor receptor (LIFR), mRNA
NM_004733	Homo sapiens acetyl-Coenzyme A transporter (ACATN), mRNA
NM_002657	Homo sapiens pleiomorphic adenoma gene-like 2 (PLAGL2), mRNA
NM_006724	Homo sapiens mitogen-activated protein kinase kinase kinase 4 (MAP3K4), transcript variant 2, mRNA
NM_006882	Homo sapiens Mdm2, transformed 3T3 cell double minute 2, p53 binding protein
	(mouse) (MDM2), transcript variant MDM2e, mRNA
NM_006881	Homo sapiens Mdm2, transformed 3T3 cell double minute 2, p53 binding protein
	(mouse) (MDM2), transcript variant MDM2d, mRNA
NM_006880	Homo sapiens Mdm2, transformed 3T3 cell double minute 2, p53 binding protein
	(mouse) (MDM2), transcript variant MDM2c, mRNA
NM_006879	Homo sapiens Mdm2, transformed 3T3 cell double minute 2, p53 binding protein
	(mouse) (MDM2), transcript variant MDM2b, mRNA
NM_006878	Homo sapiens Mdm2, transformed 3T3 cell double minute 2, p53 binding protein
	(mouse) (MDM2), transcript variant MDM2a, mRNA
NM_003801	Homo sapiens GPAA1P anchor attachment protein 1 homolog (yeast) (GPAA1), mRNA
NM_003193	Homo sapiens tubulin-specific chaperone e (TBCE), mRNA
NM_002370	Homo sapiens mago-nashi homolog, proliferation-associated (Drosophila) (MAGOH), mRNA
NM_006341	Homo sapiens MAD2 mitotic arrest deficient-like 2 (yeast) (MAD2L2), mRNA
NM_006149	Homo sapiens lectin, galactoside-binding, soluble, 4 (galectin 4) (LGALS4), mRNA
NM_003585	Homo sapiens double C2-like domains, beta (DOC2B), mRNA
NM_007129	Homo sapiens Zic family member 2 (odd-paired homolog, Drosophila) (ZIC2), mRNA
L	IIIVIA

NM 007279	Homo sapiens U2 small nuclear ribonucleoprotein auxiliary factor (65kD)
_	(U2AF65), mRNA
NM_007194	Homo sapiens CHK2 checkpoint homolog (S. pombe) (CHEK2), mRNA
NM_007271	Homo sapiens serine/threonine kinase 38 (STK38), mRNA
NM_007232	Homo sapiens histamine receptor H3 (HRH3), mRNA
NM_007278	Homo sapiens GABA(A) receptor-associated protein (GABARAP), mRNA
NM_007197	Homo sapiens frizzled homolog 10 (Drosophila) (FZD10), mRNA
NM_007246	Homo sapiens kelch-like 2, Mayven (Drosophila) (KLHL2), mRNA
NM_001466	Homo sapiens frizzled homolog 2 (Drosophila) (FZD2), mRNA
NM_006482	Homo sapiens dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 2 (DYRK2), transcript variant 2, mRNA
NM_003583	Homo sapiens dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 2 (DYRK2), transcript variant 1, mRNA
NM_006484	Homo sapiens dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 1B (DYRK1B), transcript variant c, mRNA
NM_006483	Homo sapiens dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 1B (DYRK1B), transcript variant b, mRNA
NM_001882	Homo sapiens corticotropin releasing hormone binding protein (CRHBP), mRNA
NM_005889	Homo sapiens apolipoprotein B mRNA editing enzyme, catalytic polypeptide 1 (APOBEC1), transcript variant 2, mRNA
NM_001644	Homo sapiens apolipoprotein B mRNA editing enzyme, catalytic polypeptide 1 (APOBEC1), transcript variant 1, mRNA
NM_006936	Homo sapiens SMT3 suppressor of mif two 3 homolog 1 (yeast) (SMT3H1), mRNA
NM_006912	Homo sapiens Ric-like, expressed in many tissues (Drosophila) (RIT), mRNA
NM 006910	Homo sapiens retinoblastoma binding protein 6 (RBBP6), mRNA
NM_007068	Homo sapiens DMC1 dosage suppressor of mck1 homolog, meiosis-specific homologous recombination (yeast) (DMC1), mRNA
NM 007021	Homo sapiens decidual protein induced by progesterone (DEPP), mRNA
NM_007007	Homo sapiens cleavage and polyadenylation specific factor 6, 68kD subunit (CPSF6), mRNA
NM_006822	Homo sapiens GTP-binding protein homologous to Saccharomyces cerevisiae SEC4 (SEC4L), mRNA
NM 006843	Homo sapiens serine dehydratase (SDS), mRNA
NM 006746	Homo sapiens sex comb on midleg-like 1 (Drosophila) (SCML1), mRNA
NM 006824	Homo sapiens EBNA1 binding protein 2 (EBNA1BP2), mRNA
NM_005922	Homo sapiens mitogen-activated protein kinase kinase kinase 4 (MAP3K4), transcript variant 1, mRNA
NM_006807	Homo sapiens chromobox homolog 1 (HP1 beta homolog Drosophila ) (CBX1), mRNA
NM_006734	Homo sapiens human immunodeficiency virus type I enhancer binding protein 2 (HIVEP2), mRNA
NM_006732	Homo sapiens FBJ murine osteosarcoma viral oncogene homolog B (FOSB), mRNA
NM_006729	Homo sapiens diaphanous homolog 2 (Drosophila) (DIAPH2), transcript variant 156, mRNA
NM 006829	Homo sapiens adipose specific 2 (APM2), mRNA
NM_006872	Homo sapiens TFIIA-alpha/beta-like factor (ALF), mRNA
NM_006796	Homo sapiens AFG3 ATPase family gene 3-like 2 (yeast) (AFG3L2), nuclear gene encoding mitochondrial protein, mRNA
NM 006544	Homo sapiens SEC10-like 1 (S. cerevisiae) (SEC10L1), mRNA
	,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,

NM_006666	Homo sapiens RuvB-like 2 (E. coli) (RUVBL2), mRNA
NM_006509	Homo sapiens v-rel reticuloendotheliosis viral oncogene homolog B, nuclear
	factor of kappa light polypeptide gene enhancer in B-cells 3 (avian) (RELB), mRNA
NM_006606	Homo sapiens retinoblastoma binding protein 9 (RBBP9), mRNA
NM_006620	Homo sapiens HBS1-like (S. cerevisiae) (HBS1L), mRNA
NM_006561	Homo sapiens CUG triplet repeat, RNA binding protein 2 (CUGBP2), mRNA
NM_006579	Homo sapiens emopamil binding protein (sterol isomerase) (EBP), mRNA
NM_006560	Homo sapiens CUG triplet repeat, RNA binding protein 1 (CUGBP1), mRNA
NM_001211	Homo sapiens BUB1 budding uninhibited by benzimidazoles 1 homolog beta (yeast) (BUB1B), mRNA
NM_006374	Homo sapiens serine/threonine kinase 25 (STE20 homolog, yeast) (STK25), mRNA
NM_006377	Homo sapiens unc-13-like (C. elegans) (UNC13), mRNA
NM_006357	Homo sapiens ubiquitin-conjugating enzyme E2E 3 (UBC4/5 homolog, yeast) (UBE2E3), mRNA
NM_006323	Homo sapiens SEC24 related gene family, member B (S. cerevisiae) (SEC24B), mRNA
NM_006364	Homo sapiens Sec23 homolog A (S. cerevisiae) (SEC23A), mRNA
NM_006272	Homo sapiens S100 calcium binding protein, beta (neural) (S100B), mRNA
NM_006271	Homo sapiens S100 calcium binding protein A1 (S100A1), mRNA
NM_006391	Homo sapiens RAN binding protein 7 (RANBP7), mRNA
NM_006265	Homo sapiens RAD21 homolog (S. pombe) (RAD21), mRNA
NM_006203	Homo sapiens phosphodiesterase 4D, cAMP-specific (phosphodiesterase E3 dunce homolog, Drosophila) (PDE4D), mRNA
NM_006202	Homo sapiens phosphodiesterase 4A, cAMP-specific (phosphodiesterase E2 dunce homolog, Drosophila) (PDE4A), mRNA
NM_006190	Homo sapiens origin recognition complex, subunit 2-like (yeast) (ORC2L), mRNA
NM 006181	Homo sapiens netrin 2-like (chicken) (NTN2L), mRNA
NM_006168	Homo sapiens NK6 transcription factor homolog A (Drosophila) (NKX6A), mRNA
NM_006167	Homo sapiens NK3 transcription factor homolog A (Drosophila) (NKX3A), mRNA
NM_006159	Homo sapiens NEL-like 2 (chicken) (NELL2), mRNA
NM_006157	Homo sapiens NEL-like 1 (chicken) (NELL1), mRNA
NM_005360	Homo sapiens v-maf musculoaponeurotic fibrosarcoma oncogene homolog (avian) (MAF), mRNA
NM_006306	Homo sapiens SMC1 structural maintenance of chromosomes 1-like 1 (yeast) (SMC1L1), mRNA
NM_006461	Homo sapiens mitotic spindle coiled-coil related protein (DEEPEST), mRNA
NM_006314	Homo sapiens connector enhancer of KSR-like (Drosophila kinase suppressor of ras) (CNK1), mRNA
NM_006366	Homo sapiens adenylyl cyclase-associated protein 2 (CAP2), mRNA
NM_006444	Homo sapiens SMC2 structural maintenance of chromosomes 2-like 1 (yeast) (SMC2L1), mRNA
NM_006321	Homo sapiens ariadne homolog 2 (Drosophila) (ARIH2), mRNA
NM_006406	Homo sapiens peroxiredoxin 4 (PRDX4), mRNA
NM 006334	Homo sapiens olfactomedin 1 (OLFM1), transcript variant 2, mRNA
NM 004032	Homo sapiens D-aspartate oxidase (DDO), transcript variant 2, mRNA
NM_005985	Homo sapiens snail 1 homolog, zinc finger protein (Drosophila) (SNAI1), mRNA

	The state of the s
NM_006109	Homo sapiens SKB1 homolog (S. pombe) (SKB1), mRNA
NM_005982	Homo sapiens sine oculis homeobox homolog 1 (Drosophila) (SIX1), mRNA
NM_006089	Homo sapiens sex comb on midleg-like 2 (Drosophila) (SCML2), mRNA
NM_005980	Homo sapiens S100 calcium binding protein P (S100P), mRNA
NM 005979	Homo sapiens S100 calcium binding protein A13 (S100A13), mRNA
NM 005938	Homo sapiens myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog,
_	Drosophila); translocated to, 7 (MLLT7), mRNA
NM 005937	Homo sapiens myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog,
1111_003557	Drosophila); translocated to, 6 (MLLT6), mRNA
NM_005936	Homo sapiens myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog,
11111_003550	Drosophila); translocated to, 4 (MLLT4), mRNA
NM_005935	Homo sapiens myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog,
14141_003933	Drosophila); translocated to, 2 (MLLT2), mRNA
ND4 005024	Homo sapiens myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog,
NM_005934	
) T 5 00 5000	Drosophila); translocated to, 1 (MLLT1), mRNA
NM_005933	Homo sapiens myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog,
	Drosophila) (MLL), mRNA
NM_005905	Homo sapiens MAD, mothers against decapentaplegic homolog 9 (Drosophila)
	(MADH9), mRNA
NM_005904	Homo sapiens MAD, mothers against decapentaplegic homolog 7 (Drosophila)
	(MADH7), mRNA
NM_005903	Homo sapiens MAD, mothers against decapentaplegic homolog 5 (Drosophila)
	(MADH5), mRNA
NM 005902	Homo sapiens MAD, mothers against decapentaplegic homolog 3 (Drosophila)
_	(MADH3), mRNA
NM 005901	Homo sapiens MAD, mothers against decapentaplegic homolog 2 (Drosophila)
_	(MADH2), mRNA
NM 005900	Homo sapiens MAD, mothers against decapentaplegic homolog 1 (Drosophila)
	(MADH1), mRNA
NM 006033	Homo sapiens lipase, endothelial (LIPG), mRNA
NM 006048	Homo sapiens ubiquitination factor E4B (UFD2 homolog, yeast) (UBE4B),
	mRNA
NM_006111	Homo sapiens acetyl-Coenzyme A acyltransferase 2 (mitochondrial 3-oxoacyl-
14141_000111	Coenzyme A thiolase) (ACAA2), nuclear gene encoding mitochondrial protein,
	mRNA
NM 006012	Homo sapiens ClpP caseinolytic protease, ATP-dependent, proteolytic subunit
19191_000012	homolog (E. coli) (CLPP), nuclear gene encoding mitochondrial protein, mRNA
ND ( 000110	
NM_006110	Homo sapiens CD2 antigen (cytoplasmic tail) binding protein 2 (CD2BP2),
NR 6 00 6015	mRNA
NM_006017	Homo sapiens prominin-like 1 (mouse) (PROML1), mRNA
NM_004010	Homo sapiens dystrophin (muscular dystrophy, Duchenne and Becker types),
	includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS269,
	DXS270, DXS272 (DMD), transcript variant Dp427p2, mRNA
NM_004023	Homo sapiens dystrophin (muscular dystrophy, Duchenne and Becker types),
	includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS269,
	DXS270, DXS272 (DMD), transcript variant Dp140bc, mRNA
NM_004022	Homo sapiens dystrophin (muscular dystrophy, Duchenne and Becker types),
-	includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS269,
1	DXS270, DXS272 (DMD), transcript variant D140ab, mRNA
NM 004021	Homo sapiens dystrophin (muscular dystrophy, Duchenne and Becker types),
	includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS269,
1	DXS270, DXS272 (DMD), transcript variant Dp140b, mRNA
	[

NM_004020	Homo sapiens dystrophin (muscular dystrophy, Duchenne and Becker types),
	meludes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS269,
	DXS270, DXS272 (DMD), transcript variant Dp140c, mRNA
NM_004019	Homo sapiens dystrophin (muscular dystrophy, Duchenne and Becker types).
	includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS269,
	DXS270, DXS272 (DMD), transcript variant Dp40, mRNA
NM_004018	Homo sapiens dystrophin (muscular dystrophy, Duchenne and Becker types),
	includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS269,
	DXS270, DXS272 (DMD), transcript variant Dp71ab, mRNA
NM_004017	Homo sapiens dystrophin (muscular dystrophy, Duchenne and Becker types),
Ì	includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS269,
	DXS270, DXS272 (DMD), transcript variant Dp71a, mRNA
NM_004016	Homo sapiens dystrophin (muscular dystrophy, Duchenne and Becker types),
	includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS269,
	DXS270, DXS272 (DMD), transcript variant Dp71b, mRNA
NM_004015	Homo sapiens dystrophin (muscular dystrophy, Duchenne and Becker types),
	includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS269,
	DXS270, DXS272 (DMD), transcript variant Dp71, mRNA
NM_004014	Homo sapiens dystrophin (muscular dystrophy, Duchenne and Becker types),
	includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS269,
	DXS270, DXS272 (DMD), transcript variant Dp116, mRNA
NM_004013	Homo sapiens dystrophin (muscular dystrophy, Duchenne and Becker types),
	includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS269,
	DXS270, DXS272 (DMD), transcript variant Dp140, mRNA
NM_004012	Homo sapiens dystrophin (muscular dystrophy, Duchenne and Becker types).
	includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS269
	DXS270, DXS272 (DMD), transcript variant Dp260-2, mRNA
NM_004011	Homo sapiens dystrophin (muscular dystrophy, Duchenne and Becker types)
	includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS269.
	DXS270, DXS272 (DMD), transcript variant Dp260-1, mRNA
NM_004009	Homo sapiens dystrophin (muscular dystrophy, Duchenne and Becker types),
	includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS269
\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\	DXS270, DXS272 (DMD), transcript variant Dp427p1, mRNA
NM_004007	Homo sapiens dystrophin (muscular dystrophy, Duchenne and Becker types).
	includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS269,
375 00 100 5	DXS270, DXS272 (DMD), transcript variant Dp427l, mRNA
NM_004006	Homo sapiens dystrophin (muscular dystrophy, Duchenne and Becker types),
	includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS269
NR 6 000100	DAS270, DAS272 (DMD), transcript variant Dp427m, mRNA
NM_000109	Homo sapiens dystrophin (muscular dystrophy, Duchenne and Becker types),
	includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS269,
>> 005655	DXS270, DXS272 (DMD), transcript variant Dp427c, mRNA
NM_005657	Homo sapiens tumor protein p53 binding protein, 1 (TP53BP1), mRNA
NM_005632	Homo sapiens small optic lobes homolog (Drosophila) (SOLH), mRNA
NM_005631	Homo sapiens smoothened homolog (Drosophila) (SMOH), mRNA
NM_005621	Homo sapiens \$100 calcium binding protein A12 (calgranulin C) (\$100A12).
ND 6 005500	mRNA
NM_005620	Homo sapiens S100 calcium binding protein A11 (calgizzarin) (S100A11),
NB ( 007610	mRNA
NM_005610	Homo sapiens retinoblastoma binding protein 4 (RBBP4), mRNA
NM_005732	Homo sapiens RAD50 homolog (S. cerevisiae) (RAD50), mRNA
NM_005591	Homo sapiens MRE11 meiotic recombination 11 homolog A (S. cerevisiae)
	(MRE11A), mRNA

NM_005590	Homo sapiens MRE11 meiotic recombination 11 homolog A (S. cerevisiae) (MRE11A), mRNA
NM 005585	Homo sapiens MAD, mothers against decapentaplegic homolog 6 (Drosophila)
1414_005585	(MADH6), mRNA
NM 005584	Homo sapiens mab-21-like 1 (C. elegans) (MAB21L1), mRNA
NM 005582	Homo sapiens lymphocyte antigen 64 homolog, radioprotective 105kD (mouse)
11	(LY64), mRNA
NM 005667	Homo sapiens zinc finger protein 103 homolog (mouse) (ZFP103), mRNA
NM 005886	Homo sapiens katanin p80 (WD40-containing) subunit B 1 (KATNB1), mRNA
NM 005860	Homo sapiens follistatin-like 3 (secreted glycoprotein) (FSTL3), mRNA
NM 005758	Homo sapiens heterogeneous nuclear ribonucleoprotein A3 (HNRPA3), mRNA
NM 005510	Homo sapiens dom-3 homolog Z (C. elegans) (DOM3Z), transcript variant 2,
	mRNA
NM_005766	Homo sapiens FERM, RhoGEF (ARHGEF) and pleckstrin domain protein 1
	(chondrocyte-derived) (FARP1), mRNA
NM 005722	Homo sapiens ARP2 actin-related protein 2 homolog (yeast) (ACTR2), mRNA
NM 005750	Homo sapiens chromosome 4 open reading frame 6 (C4orf6), mRNA
NM 005170	Homo sapiens achaete-scute complex-like 2 (Drosophila) (ASCL2), mRNA
NM 005426	Homo sapiens tumor protein p53 binding protein, 2 (TP53BP2), mRNA
NM 005486	Homo sapiens target of myb1-like 1 (chicken) (TOM1L1), mRNA
NM 005488	Homo sapiens target of myb1 (chicken) (TOM1), mRNA
NM 005417	Homo sapiens v-src sarcoma (Schmidt-Ruppin A-2) viral oncogene homolog
NWI_003417	(avian) (SRC), mRNA
NTM 005412	Homo sapiens sine oculis homeobox homolog 3 (Drosophila) (SIX3), mRNA
NM_005413	
NM_005444	Homo sapiens RCD1 required for cell differentiation1 homolog (S. pombe) (RQCD1), mRNA
NM_005378	Homo sapiens v-myc myelocytomatosis viral related oncogene, neuroblastoma derived (avian) (MYCN), mRNA
NM_005377	Homo sapiens v-myc myelocytomatosis viral oncogene homolog 2 (avian) (MYCL2), mRNA
NM_005375	Homo sapiens v-myb myeloblastosis viral oncogene homolog (avian) (MYB), mRNA
NM_005359	Homo sapiens MAD, mothers against decapentaplegic homolog 4 (Drosophila) (MADH4), mRNA
NM 005340	Homo sapiens histidine triad nucleotide binding protein (HINT), mRNA
NM_005307	Homo sapiens G protein-coupled receptor kinase 2-like (Drosophila) (GPRK2L), mRNA
NM_005262	Homo sapiens growth factor, augmenter of liver regeneration (ERV1 homolog, S. cerevisiae) (GFER), mRNA
NM_005261	Homo sapiens GTP binding protein overexpressed in skeletal muscle (GEM), mRNA
NM 005257	Homo sapiens GATA binding protein 6 (GATA6), mRNA
NM 005245	Homo sapiens FAT tumor suppressor homolog 1 (Drosophila) (FAT), mRNA
NM 005244	Homo sapiens eyes absent homolog 2 (Drosophila) (EYA2), mRNA
NM 005239	Homo sapiens v-ets erythroblastosis virus E26 oncogene homolog 2 (avian)
14141_003239	(ETS2), mRNA
NM_005235	Homo sapiens v-erb-a erythroblastic leukemia viral oncogene homolog 4 (avian) (ERBB4), mRNA
NM_005228	Homo sapiens epidermal growth factor receptor (erythroblastic leukemia viral (verb-b) oncogene homolog, avian) (EGFR), mRNA
NM 005224	Homo sapiens dead ringer-like 1 (Drosophila) (DRIL1), mRNA
NM 005219	Homo sapiens diaphanous homolog 1 (Drosophila) (DIAPH1), mRNA
11112 300217	1 Arabamana managa - (

Nom   Nome   N		
NM_005197   Homo sapiens checkpoint suppressor 1 (CHES1), mRNA	NM_005207	Homo sapiens v-crk sarcoma virus CT10 oncogene homolog (avian)-like (CRKL), mRNA
NM_005454   Homo sapiens cerberus 1 homolog, cysteine knot superfamily (Xenopus laevis) (CER1), mRNA   NM_005466   Homo sapiens SMC4 structural maintenance of chromosomes 4-like 1 (yeast) (SMC4L1), mRNA   Homo sapiens sistaless homeobox (Drosophila) (ARIX), mRNA   NM_005078   Homo sapiens transducin-like enhancer of split 3 (E(sp1) homolog, Drosophila) (TLE1), mRNA   Homo sapiens transducin-like enhancer of split 1 (E(sp1) homolog, Drosophila) (TLE1), mRNA   Homo sapiens single-minded homolog 1 (Drosophila) (SIM1), mRNA   NM_005067   Homo sapiens single-minded homolog 2 (Drosophila) (SIM1), mRNA   NM_005067   Homo sapiens seven in absentia homolog 2 (Drosophila) (SIAH2), mRNA   NM_005138   Homo sapiens SCO cytochrome oxidase deficient homolog 2 (yeast) (SCO2), nuclear gene encoding mitochondrial protein, mRNA   NM_005136   Homo sapiens RCE1 homolog, prenyl protein protease (S. cerevisiae) (RCE1), mRNA   NM_005056   Homo sapiens retinoblastoma binding protein 2 (RBBP5), mRNA   NM_005056   Homo sapiens retinoblastoma binding protein 2 (RBBP5), mRNA   NM_005056   Homo sapiens RAD23 homolog A (S. cerevisiae) (RAD23A), mRNA   NM_005056   Homo sapiens PWP2 periodic tryptophan protein homolog (yeast) (PWP2H), mRNA   NM_005066   Homo sapiens PWP2 periodic tryptophan protein homolog (yeast) (PWP2H), mRNA   NM_005066   Homo sapiens retinoblastoma binding protein 2 (RBBP2), mRNA   NM_005066   Homo sapiens retinoblastoma binding protein 2 (RBBP3), mRNA   NM_005066   Homo sapiens SWP2 periodic tryptophan protein homolog (yeast) (PWP2H), mRNA   NM_00467   Homo sapiens binding protein H (MYBPH), mRNA   NM_00467   Homo sapiens binding protein H (MYBPH), mRNA   NM_00467   Homo sapiens transmembrane 4 superfamily member 4 (TM4SF4), mRNA   NM_00467   Homo sapiens staufen, RNA binding protein (Drosophila) (STAU), transcript variant T4, mRNA   Homo sapiens staufen, RNA binding protein (Drosophila) (STAU), transcript variant T4, mRNA   Homo sapiens staufen, RNA binding protein (Drosophila) (STAU), transcript variant 1, mRNA   Homo	NM 005197	
NM_005169   Homo sapiens SMC4 structural maintenance of chromosomes 4-like 1 (yeast) (SMC4L1), mRNA   NM_005169   Homo sapiens aristaless homeobox (Drosophila) (ARIX), mRNA   NM_005078   Homo sapiens transducin-like enhancer of split 3 (E(sp1) homolog, Drosophila) (TLB3), mRNA   NM_005077   Homo sapiens transducin-like enhancer of split 1 (E(sp1) homolog, Drosophila) (TLB1), mRNA   NM_005068   Homo sapiens single-minded homolog 1 (Drosophila) (SIM1), mRNA   NM_005067   Homo sapiens single-minded homolog 2 (Drosophila) (SIM12), mRNA   NM_005138   Homo sapiens seven in absentia homolog 2 (Drosophila) (SIAH2), mRNA   NM_005164   Homo sapiens ROD1 regulator of differentiation 1 (S, pombe) (ROD1), mRNA   NM_005156   Homo sapiens RCD1 homolog, prenyl protein protease (S. cerevisiae) (RCE1), mRNA   NM_005057   Homo sapiens retinoblastoma binding protein 5 (RBBP5), mRNA   NM_005056   Homo sapiens retinoblastoma binding protein 2 (RBBP5), mRNA   NM_005056   Homo sapiens retinoblastoma binding protein 2 (RBBP5), mRNA   NM_005056   Homo sapiens RAD23 homolog A (S. cerevisiae) (RAD23A), mRNA   NM_005056   Homo sapiens PWP2 periodic tryptophan protein homolog (yeast) (PWP2H), mRNA   NM_005060   Homo sapiens PWP2 periodic tryptophan protein 1 (RBBP5), mRNA   NM_005060   Homo sapiens myosin binding protein H (MYBPH), mRNA   NM_004607   Homo sapiens myosin binding protein H (MYBPH), mRNA   NM_004677   Homo sapiens transmembrane 4 superfamily member 4 (TM4SF4), mRNA   NM_004677   Homo sapiens transmembrane 4 superfamily member 4 (TM4SF4), mRNA   NM_004678   Homo sapiens strundenance   Homosome   NM_004679   Homo sapiens studin-specific chaperone a (TBCA), mRNA   NM_004679   Homo sapiens studin-specific chaperone a (TBCA), mRNA   NM_004674   Homo sapiens studin-specific paperone (mouse) (SMCY), mRNA   NM_004674   Homo sapiens studin-specific paperone (mouse) (SMCY), mRNA   NM_004674   Homo sapiens sevenum deprivation response (phosphatidylserine binding protein) (SDPR), mRNA   NM_004584   Homo sapiens retinol binding protein 1	NM_005454	Homo sapiens cerberus 1 homolog, cysteine knot superfamily (Xenopus laevis)
NM_005078   Homo sapiens aristaless homeobox (Drosophila) (ARIX), mRNA   NM_005078   Homo sapiens transducin-like enhancer of split 3 (E(spl1) homolog, Drosophila) (TLE3), mRNA   NM_005067   Homo sapiens transducin-like enhancer of split 1 (E(spl1) homolog, Drosophila) (TLE1), mRNA   Homo sapiens single-minded homolog 1 (Drosophila) (SIM1), mRNA   NM_005067   Homo sapiens seven in absentia homolog 2 (Drosophila) (SIAH2), mRNA   NM_005138   Homo sapiens SCO cytochrome oxidase deficient homolog 2 (yeast) (SCO2), nuclear gene encoding mitochondrial protein, mRNA   NM_005156   Homo sapiens RCD1 regulator of differentiation 1 (S. pombe) (ROD1), mRNA   NM_005131   Homo sapiens RCE1 homolog, prenyl protein protease (S. cerevisiae) (RCE1), mRNA   NM_005057   Homo sapiens retinoblastoma binding protein 5 (RBBF5), mRNA   NM_005056   Homo sapiens retinoblastoma binding protein 5 (RBBF5), mRNA   NM_005056   Homo sapiens RAD23 homolog A (S. cerevisiae) (RAD23A), mRNA   NM_005054   Homo sapiens RAD23 homolog A (S. cerevisiae) (RAD23A), mRNA   NM_005056   Homo sapiens RAD23 homolog A (S. cerevisiae) (RAD23A), mRNA   NM_005056   Homo sapiens NHP2 non-histone chromosome protein 2-like 1 (S. cerevisiae) (NHP2L1), mRNA   NM_005056   Homo sapiens myosin binding protein H (MYBPH), mRNA   NM_004997   Homo sapiens myosin binding protein H (MYBPH), mRNA   NM_004677   Homo sapiens Testis-specific XK-related protein on Y (XKRY), mRNA   NM_004788   Homo sapiens transmembrane 4 superfamily member 4 (TM4SF4), mRNA   NM_004671   Homo sapiens transmembrane 4 superfamily member 4 (TM4SF4), mRNA   NM_00460   Homo sapiens stubulin-specific chaperone a (TBCA), mRNA   Homo sapiens stubulin-specific chaperone a (TBCA), mRNA   Homo sapiens stubulin-specific chaperone a (TBCA), mRNA   Homo sapiens selit homolog 2 (Drosophila) (SLIT2), mRNA   Homo sapiens selic trafficking protein (SEC22C), transcript variant 2, mRNA   Homo sapiens selic trafficking protein (SEC22C), transcript variant 2, mRNA   Homo sapiens select trafficking protein (SEC22C), tra	NM_005496	Homo sapiens SMC4 structural maintenance of chromosomes 4-like 1 (yeast)
NM_005078  Homo sapiens transducin-like enhancer of split 3 (E(sp1) homolog, Drosophila) (TLE3), mRNA  NM_005077  Homo sapiens transducin-like enhancer of split 1 (E(sp1) homolog, Drosophila) (TLE1), mRNA  NM_005068  Homo sapiens single-minded homolog 1 (Drosophila) (SIM1), mRNA  NM_005067  Homo sapiens seven in absentia homolog 2 (Drosophila) (SIM1), mRNA  NM_005138  Homo sapiens SCO cytochrome oxidase deficient homolog 2 (yeast) (SCO2), nuclear gene encoding mitochondrial protein, mRNA  NM_005156  Homo sapiens RCE1 homolog, prenyl protein protease (S. cerevisiae) (RCE1), mRNA  NM_005157  Homo sapiens RCE1 homolog, prenyl protein protease (S. cerevisiae) (RCE1), mRNA  NM_005057  Homo sapiens retinoblastoma binding protein 2 (RBBP5), mRNA  NM_005053  Homo sapiens RD23 homolog A (S. cerevisiae) (RAD23A), mRNA  NM_005054  Homo sapiens RD23 homolog A (S. cerevisiae) (RAD23A), mRNA  NM_005055  Homo sapiens NHP2 pon-histone chromosome protein 2-like 1 (S. cerevisiae) (NHP2L1), mRNA  NM_005069  Homo sapiens single-minded homolog (yeast) (PWP2H), mRNA  NM_004697  Homo sapiens single-minded homolog (yeast) (PWP2H), mRNA  NM_004677  Homo sapiens single-minded homolog (yeast) (PWP2H), mRNA  NM_004677  Homo sapiens single-minded homolog (yeast) (WBP4A), mRNA  NM_004678  Homo sapiens single-minded homolog (yeast) (UBE4A), mRNA  NM_004674  Homo sapiens single-minded homolog (yeast) (UBE4A), mRNA  NM_004675  Homo sapiens single-minded homolog (yeast) (UBE4A), mRNA  NM_004676  Homo sapiens single-minded homolog (yeast) (UBE4A), mRNA  NM_004677  Homo sapiens single-minded homolog (yeast) (VBE4A), mRNA  NM_004678  Homo sapiens single-minded homolog (yeast) (VBE4A), mRNA  NM_004674  Homo sapiens single-minded homolog (yeast) (VBE4A), mRNA  NM_004675  Homo sapiens single-minded homolog (yeast)	NM 005169	
(TLE3), mRNA  NM_005068  Homo sapiens single-minded homolog 1 (Drosophila) (SIM1), mRNA  NM_005068  Homo sapiens single-minded homolog 1 (Drosophila) (SIM1), mRNA  NM_005067  Homo sapiens seven in absentia homolog 2 (Drosophila) (SIAH2), mRNA  NM_005138  Homo sapiens SCO cytochrome oxidase deficient homolog 2 (yeast) (SCO2), nuclear gene encoding mitochondrial protein, mRNA  NM_005156  Homo sapiens RCE1 homolog, prenyl protein protease (S. cerevisiae) (RCE1), mRNA  NM_005133  Homo sapiens RCE1 homolog, prenyl protein protease (S. cerevisiae) (RCE1), mRNA  NM_005057  Homo sapiens retinoblastoma binding protein 5 (RBBP5), mRNA  NM_005058  Homo sapiens retinoblastoma binding protein 2 (RBBP2), mRNA  NM_005049  Homo sapiens PWP2 periodic tryptophan protein homolog (yeast) (PWP2H), mRNA  NM_00508  Homo sapiens NHP2 non-histone chromosome protein 2-like 1 (S. cerevisiae) (NHP2L1), mRNA  NM_004077  Homo sapiens myosin binding protein H (MYBPH), mRNA  NM_004677  Homo sapiens setisis-specific XK-related protein on Y (XKRY), mRNA  NM_004678  Homo sapiens trusinsmembrane 4 superfamily member 4 (TM4SF4), mRNA  NM_004671  Homo sapiens tubulin-specific chaperone a (TBCA), mRNA  NM_004602  Homo sapiens Staufen, RNA binding protein (Drosophila) (STAU), transcript variant T4, mRNA  NM_004653  Homo sapiens Secure homolog, Y chromosome (mouse) (SMCY), mRNA  NM_004667  Homo sapiens Secure homolog, Y chromosome (mouse) (SMCY), mRNA  NM_004677  Homo sapiens Secure homolog, Y chromosome (mouse) (SMCY), mRNA  NM_004667  Homo sapiens Secure homolog, Y chromosome (mouse) (SMCY), mRNA  NM_004678  Homo sapiens studen, RNA binding protein (Drosophila) (STAU), transcript variant T4, mRNA  NM_004678  Homo sapiens secure deprivation response (phosphatidylserine binding protein) (SDPR), mRNA  NM_004593  Homo sapiens secure deprivation response (phosphatidylserine binding protein) (SDPR), mRNA  NM_004584  Homo sapiens RAB33A, member RAS oncogene family (RAB33A), mRNA  NM_004584  Homo sapiens peroxisomal biogenesis factor 16 (PEX16), transcript		Homo sapiens transducin-like enhancer of split 3 (F(spl) homolog Drosophila)
NM_005077	_	(TLE3), mRNA
NM_005067 NM_005138 NM_005138 Homo sapiens seven in absentia homolog 2 (Drosophila) (SIAH2), mRNA NM_005156 Homo sapiens ROD1 regulator of differentiation 1 (S. pombe) (ROD1), mRNA NM_005133 NM_005133 NM_005134 NM_005057 Homo sapiens RCE1 homolog, prenyl protein protease (S. cerevisiae) (RCE1), mRNA NM_005057 NM_005056 NM_005056 NM_005056 NM_005056 NM_005050 NM_005059 NM_005059 Homo sapiens retinoblastoma binding protein 2 (RBBP2), mRNA NM_005050 NM_00	NM_005077	Homo sapiens transducin-like enhancer of split 1 (E(sp1) homolog, Drosophila) (TLE1), mRNA
NM_005067 NM_005138 NM_005138 Homo sapiens seven in absentia homolog 2 (Drosophila) (SIAH2), mRNA NM_005156 Homo sapiens ROD1 regulator of differentiation 1 (S. pombe) (ROD1), mRNA NM_005133 NM_005133 NM_005134 NM_005057 Homo sapiens RCE1 homolog, prenyl protein protease (S. cerevisiae) (RCE1), mRNA NM_005057 NM_005056 NM_005056 NM_005056 NM_005056 NM_005050 NM_005059 NM_005059 Homo sapiens retinoblastoma binding protein 2 (RBBP2), mRNA NM_005050 NM_00	NM_005068	Homo sapiens single-minded homolog 1 (Drosophila) (SIM1), mRNA
NM_005138 Homo sapiens SCO cytochrome oxidase deficient homolog 2 (yeast) (SCO2), nuclear gene encoding mitochondrial protein, mRNA NM_005156 Homo sapiens ROD1 regulator of differentiation 1 (S. pombe) (ROD1), mRNA NM_005133 Homo sapiens RCE1 homolog, prenyl protein protease (S. cerevisiae) (RCE1), mRNA NM_005057 Homo sapiens retinoblastoma binding protein 2 (RBBP5), mRNA NM_005058 Homo sapiens RAD23 homolog A (S. cerevisiae) (RAD23A), mRNA NM_005049 Homo sapiens RAD23 homolog A (S. cerevisiae) (RAD23A), mRNA NM_005049 Homo sapiens RAD23 homolog A (S. cerevisiae) (RAD23A), mRNA NM_005008 Homo sapiens PWP2 periodic tryptophan protein homolog (yeast) (PWP2H), mRNA NM_005008 Homo sapiens nylP2 non-histone chromosome protein 2-like 1 (S. cerevisiae) (NHP2L1), mRNA NM_004677 Homo sapiens myosin binding protein H (MYBPH), mRNA NM_004677 Homo sapiens binduitination factor E4A (UFD2 homolog, yeast) (UBE4A), mRNA NM_004671 Homo sapiens transmembrane 4 superfamily member 4 (TM4SF4), mRNA NM_004607 Homo sapiens tubulin-specific chaperone a (TBCA), mRNA NM_004607 Homo sapiens staufen, RNA binding protein (Drosophila) (STAU), transcript variant T4, mRNA NM_00463 Homo sapiens Smcy homolog, Y chromosome (mouse) (SMCY), mRNA NM_004593 Homo sapiens splicing factor, arginine/serine-rich 10 (transformer 2 homolog, Drosophila) (SFRS10), mRNA NM_004594 Homo sapiens vesicle trafficking protein (SEC2CC), transcript variant 2, mRNA NM_004587 Homo sapiens serum deprivation response (phosphatidylserine binding protein) (SDPR), mRNA NM_004587 Homo sapiens ribosome binding protein (SEC2CC), transcript variant 2, mRNA NM_004589 Homo sapiens ribosome binding protein (REP2), mRNA NM_004584 Homo sapiens RAD9 homolog (S. pombe) (RAD9), mRNA NM_004584 Homo sapiens retinol binding protein 2, cellular (RBP2), mRNA NM_004584 Homo sapiens peroxisomal biogenesis factor 16 (PEX16), transcript variant 1, mRNA NM_004584 Homo sapiens peroxisomal biogenesis factor 16 (PEX16), transcript variant 1, mRNA	NM_005067	Homo sapiens seven in absentia homolog 2 (Drosophila) (SIAH2), mRNA
nuclear gene encoding mitochondrial protein, mRNA NM_005156 Homo sapiens ROD1 regulator of differentiation 1 (S. pombe) (ROD1), mRNA NM_005131 Homo sapiens RCE1 homolog, prenyl protein protease (S. cerevisiae) (RCE1), mRNA NM_005056 Homo sapiens retinoblastoma binding protein 5 (RBBP5), mRNA NM_005056 Homo sapiens retinoblastoma binding protein 2 (RBBP2), mRNA NM_005051 Homo sapiens RAD23 homolog A (S. cerevisiae) (RAD23A), mRNA NM_005053 Homo sapiens RAD23 homolog A (S. cerevisiae) (RAD23A), mRNA NM_005054 Homo sapiens PWP2 periodic tryptophan protein homolog (yeast) (PWP2H), mRNA NM_005008 Homo sapiens NHP2 non-histone chromosome protein 2-like 1 (S. cerevisiae) (NHP2L1), mRNA NM_004997 Homo sapiens myosin binding protein H (MYBPH), mRNA NM_004677 Homo sapiens stesis-specific XK-related protein on Y (XKRY), mRNA NM_00478 Homo sapiens ubiquitination factor E4A (UFD2 homolog, yeast) (UBE4A), mRNA NM_004601 Homo sapiens transmembrane 4 superfamily member 4 (TM4SF4), mRNA NM_004602 Homo sapiens staufen, RNA binding protein (Drosophila) (STAU), transcript variant T4, mRNA NM_004653 Homo sapiens Smcy homolog, Y chromosome (mouse) (SMCY), mRNA NM_004787 Homo sapiens Smcy homolog, Y chromosome (mouse) (SMCY), mRNA NM_004593 Homo sapiens splicing factor, arginine/serine-rich 10 (transformer 2 homolog, Drosophila) (SFRS10), mRNA NM_004594 Homo sapiens serum deprivation response (phosphatidylserine binding protein) (SDPR), mRNA NM_004587 Homo sapiens retinol binding protein (SEC22C), transcript variant 2, mRNA NM_004587 Homo sapiens retinol binding protein 1 homolog 1 (yeast) (SCO1), nuclear gene encoding mitochondrial protein, mRNA NM_004587 Homo sapiens retinol binding protein 2, cellular (RBP2), mRNA NM_004588 Homo sapiens RAB33A, member RAS oncogene family (RAB33A), mRNA NM_004564 Homo sapiens PET112-like (yeast) (PET112L), mRNA	NM_005138	Homo sapiens SCO cytochrome oxidase deficient homolog 2 (yeast) (SCO2),
NM005156Homo sapiens ROD1 regulator of differentiation 1 (S. pombe) (ROD1), mRNANM005133Homo sapiens RCE1 homolog, prenyl protein protease (S. cerevisiae) (RCE1), mRNANM005057Homo sapiens retinoblastoma binding protein 5 (RBBP5), mRNANM005056Homo sapiens RAD23 homolog A (S. cerevisiae) (RAD23A), mRNANM005053Homo sapiens RAD23 homolog A (S. cerevisiae) (RAD23A), mRNANM005049Homo sapiens PWP2 periodic tryptophan protein homolog (yeast) (PWP2H), mRNANM005008Homo sapiens NHP2 non-histone chromosome protein 2-like 1 (S. cerevisiae) (NHP2L1), mRNANM004997Homo sapiens myosin binding protein H (MYBPH), mRNANM004677Homo sapiens Testis-specific XK-related protein on Y (XKRY), mRNANM004678Homo sapiens transmembrane 4 superfamily member 4 (TM4SF4), mRNANM004607Homo sapiens transmembrane 4 superfamily member 4 (TM4SF4), mRNANM004607Homo sapiens staufen, RNA binding protein (Drosophila) (STAU), transcript variant T4, mRNANM00463Homo sapiens Smcy homolog, Y chromosome (mouse) (SMCY), mRNANM00453Homo sapiens slit homolog 2 (Drosophila) (SLIT2), mRNANM004593Homo sapiens slit fomolog 2 (Drosophila) (SLT2), mRNANM004567Homo sapiens serum deprivation response (phosphatidylserine binding protein) (SDPR), mRNANM004589Homo sapiens serum deprivation response (phosphatidylserine binding protein) (SDPR), mRNANM004584Homo sapiens retinol binding protein 1, mRNANM004584		nuclear gene encoding mitochondrial protein, mRNA
NM_00505133   Homo sapiens RCE1 homolog, prenyl protein protease (S. cerevisiae) (RCE1), mRNA   NM_005055   Homo sapiens retinoblastoma binding protein 5 (RBBP5), mRNA   NM_005056   Homo sapiens RAD23 homolog A (S. cerevisiae) (RAD23A), mRNA   Homo sapiens PWP2 periodic tryptophan protein homolog (yeast) (PWP2H), mRNA   NM_005049   Homo sapiens PWP2 periodic tryptophan protein homolog (yeast) (PWP2H), mRNA   NM_005008   Homo sapiens NHP2 non-histone chromosome protein 2-like 1 (S. cerevisiae) (NHP2L1), mRNA   Homo sapiens myosin binding protein H (MYBPH), mRNA   NM_004677   Homo sapiens Testis-specific XK-related protein on Y (XKRY), mRNA   NM_004678   Homo sapiens transmembrane 4 superfamily member 4 (TM4SF4), mRNA   NM_004601   Homo sapiens studilin-specific chaperone a (TBCA), mRNA   Homo sapiens smcy homolog, Y chromosome (mouse) (SMCY), mRNA   NM_004602   Homo sapiens slit homolog 2 (Drosophila) (SITAU), transcript variant T4, mRNA   Homo sapiens splicing factor, arginine/serine-rich 10 (transformer 2 homolog, Drosophila) (SFRS10), mRNA   Homo sapiens serum deprivation response (phosphatidylserine binding protein) (SDPR), mRNA   Homo sapiens srum deprivation response (phosphatidylserine binding protein) (SDPR), mRNA   Homo sapiens retinol binding protein 1 homolog 1 (yeast) (SCO1), nuclear gene encoding mitochondrial protein, mRNA   Homo sapiens retinol binding protein 2, cellular (RBP2), mRNA   Homo sapiens RAD9 homolog (S. pombe) (RAD9), mRNA   Homo sapiens RAD9 homolog (S. pombe) (RAD9), mRNA   Homo sapiens PET112-like (yeast) (PET112L), mRNA   Homo sapiens PET112-like (yeast) (PE	NM_005156	Homo sapiens ROD1 regulator of differentiation 1 (S. pombe) (ROD1), mRNA
NM_005055   Homo sapiens retinoblastoma binding protein 2 (RBBP2), mRNA	NM_005133	Homo sapiens RCE1 homolog, prenyl protein protease (S. cerevisiae) (RCE1),
NM_005055   Homo sapiens retinoblastoma binding protein 2 (RBBP2), mRNA	NM 005057	Homo sapiens retinoblastoma binding protein 5 (RBBP5), mRNA
NM 005053   Homo sapiens RAD23 homolog A (S. cerevisiae) (RAD23A), mRNA	NM 005056	Homo sapiens retinoblastoma binding protein 2 (RBBP2), mRNA
NM_005049   Homo sapiens PWP2 periodic tryptophan protein homolog (yeast) (PWP2H), mRNA	NM 005053	Homo sapiens RAD23 homolog A (S. cerevisiae) (RAD23A), mRNA
NM_004997		Homo sapiens PWP2 periodic tryptophan protein homolog (yeast) (PWP2H),
NM_004997 Homo sapiens myosin binding protein H (MYBPH), mRNA NM_004677 Homo sapiens Testis-specific XK-related protein on Y (XKRY), mRNA NM_004788 Homo sapiens ubiquitination factor E4A (UFD2 homolog, yeast) (UBE4A), mRNA NM_004617 Homo sapiens transmembrane 4 superfamily member 4 (TM4SF4), mRNA NM_004607 Homo sapiens tubulin-specific chaperone a (TBCA), mRNA NM_004602 Homo sapiens staufen, RNA binding protein (Drosophila) (STAU), transcript variant T4, mRNA NM_004653 Homo sapiens Smcy homolog, Y chromosome (mouse) (SMCY), mRNA NM_004787 Homo sapiens slit homolog 2 (Drosophila) (SLIT2), mRNA NM_004593 Homo sapiens splicing factor, arginine/serine-rich 10 (transformer 2 homolog, Drosophila) (SFRS10), mRNA NM_004657 Homo sapiens vesicle trafficking protein (SEC22C), transcript variant 2, mRNA NM_004657 Homo sapiens serum deprivation response (phosphatidylserine binding protein) (SDPR), mRNA NM_004589 Homo sapiens SCO cytochrome oxidase deficient homolog 1 (yeast) (SCO1), nuclear gene encoding mitochondrial protein, mRNA NM_004587 Homo sapiens retinol binding protein 1 homolog 180kD (dog) (RRBP1), mRNA NM_004584 Homo sapiens RAD9 homolog (S. pombe) (RAD9), mRNA NM_004794 Homo sapiens RAB33A, member RAS oncogene family (RAB33A), mRNA NM_004564 Homo sapiens PET112-like (yeast) (PET112L), mRNA	NM_005008	Homo sapiens NHP2 non-histone chromosome protein 2-like 1 (S. cerevisiae)
NM_004677 Homo sapiens Testis-specific XK-related protein on Y (XKRY), mRNA NM_004788 Homo sapiens ubiquitination factor E4A (UFD2 homolog, yeast) (UBE4A), mRNA NM_004617 Homo sapiens transmembrane 4 superfamily member 4 (TM4SF4), mRNA NM_004607 Homo sapiens tubulin-specific chaperone a (TBCA), mRNA NM_004602 Homo sapiens staufen, RNA binding protein (Drosophila) (STAU), transcript variant T4, mRNA NM_004653 Homo sapiens Smcy homolog, Y chromosome (mouse) (SMCY), mRNA NM_004787 Homo sapiens slit homolog 2 (Drosophila) (SLIT2), mRNA NM_004593 Homo sapiens splicing factor, argimine/serine-rich 10 (transformer 2 homolog, Drosophila) (SFRS10), mRNA NM_004657 Homo sapiens vesicle trafficking protein (SEC22C), transcript variant 2, mRNA NM_004658 Homo sapiens serum deprivation response (phosphatidylserine binding protein) (SDPR), mRNA NM_004589 Homo sapiens SCO cytochrome oxidase deficient homolog 1 (yeast) (SC01), nuclear gene encoding mitochondrial protein, mRNA NM_004587 Homo sapiens retinol binding protein 2, cellular (RBP2), mRNA NM_004584 Homo sapiens RAD9 homolog (S. pombe) (RAD9), mRNA NM_004584 Homo sapiens RAD9 homolog (S. pombe) (RAD9), mRNA NM_004584 Homo sapiens RAB33A, member RAS oncogene family (RAB33A), mRNA NM_004564 Homo sapiens PET112-like (yeast) (PET112L), mRNA	NM 004997	
NM_004617 Homo sapiens ubiquitination factor E4A (UFD2 homolog, yeast) (UBE4A), mRNA  NM_004607 Homo sapiens transmembrane 4 superfamily member 4 (TM4SF4), mRNA  NM_004602 Homo sapiens tubulin-specific chaperone a (TBCA), mRNA  NM_004602 Homo sapiens staufen, RNA binding protein (Drosophila) (STAU), transcript variant T4, mRNA  NM_004653 Homo sapiens Smcy homolog, Y chromosome (mouse) (SMCY), mRNA  NM_004787 Homo sapiens splicing factor, arginine/serine-rich 10 (transformer 2 homolog, Drosophila) (SFRS10), mRNA  NM_004593 Homo sapiens vesicle trafficking protein (SEC22C), transcript variant 2, mRNA  NM_004206 Homo sapiens vesicle trafficking protein (SEC22C), transcript variant 2, mRNA  NM_004567 Homo sapiens serum deprivation response (phosphatidylserine binding protein) (SDPR), mRNA  NM_004589 Homo sapiens SCO cytochrome oxidase deficient homolog 1 (yeast) (SCO1), nuclear gene encoding mitochondrial protein, mRNA  NM_004587 Homo sapiens retinol binding protein 1 homolog 180kD (dog) (RRBP1), mRNA  NM_004584 Homo sapiens RAD9 homolog (S. pombe) (RAD9), mRNA  NM_004584 Homo sapiens RAB33A, member RAS oncogene family (RAB33A), mRNA  NM_004813 Homo sapiens PET112-like (yeast) (PET112L), mRNA	NM 004677	Homo sapiens Testis-specific XK-related protein on Y (XKRY), mRNA
NM_004607 Homo sapiens tubulin-specific chaperone a (TBCA), mRNA  NM_004602 Homo sapiens staufen, RNA binding protein (Drosophila) (STAU), transcript variant T4, mRNA  NM_004653 Homo sapiens Smcy homolog, Y chromosome (mouse) (SMCY), mRNA  NM_004787 Homo sapiens slit homolog 2 (Drosophila) (SLTT2), mRNA  NM_004593 Homo sapiens splicing factor, arginine/serine-rich 10 (transformer 2 homolog, Drosophila) (SFRS10), mRNA  NM_004206 Homo sapiens vesicle trafficking protein (SEC22C), transcript variant 2, mRNA  NM_004657 Homo sapiens serum deprivation response (phosphatidylserine binding protein) (SDPR), mRNA  NM_004589 Homo sapiens SCO cytochrome oxidase deficient homolog 1 (yeast) (SCO1), nuclear gene encoding mitochondrial protein, mRNA  NM_004587 Homo sapiens ribosome binding protein 1 homolog 180kD (dog) (RRBP1), mRNA  NM_004164 Homo sapiens retinol binding protein 2, cellular (RBP2), mRNA  NM_004584 Homo sapiens RAD9 homolog (S. pombe) (RAD9), mRNA  NM_004794 Homo sapiens RAB33A, member RAS oncogene family (RAB33A), mRNA  NM_004813 Homo sapiens peroxisomal biogenesis factor 16 (PEX16), transcript variant 1, mRNA  NM_004564 Homo sapiens PET112-like (yeast) (PET112L), mRNA	NM_004788	Homo sapiens ubiquitination factor E4A (UFD2 homolog, yeast) (UBE4A),
NM_004607 Homo sapiens tubulin-specific chaperone a (TBCA), mRNA  NM_004602 Homo sapiens staufen, RNA binding protein (Drosophila) (STAU), transcript variant T4, mRNA  NM_004653 Homo sapiens Smcy homolog, Y chromosome (mouse) (SMCY), mRNA  NM_004787 Homo sapiens slit homolog 2 (Drosophila) (SLIT2), mRNA  NM_004593 Homo sapiens splicing factor, arginine/serine-rich 10 (transformer 2 homolog, Drosophila) (SFRS10), mRNA  NM_004206 Homo sapiens vesicle trafficking protein (SEC22C), transcript variant 2, mRNA  NM_004567 Homo sapiens serum deprivation response (phosphatidylserine binding protein) (SDPR), mRNA  NM_004589 Homo sapiens SCO cytochrome oxidase deficient homolog 1 (yeast) (SCO1), nuclear gene encoding mitochondrial protein, mRNA  NM_004587 Homo sapiens retinol binding protein 2, cellular (RBP2), mRNA  NM_004584 Homo sapiens RAD9 homolog (S. pombe) (RAD9), mRNA  NM_004794 Homo sapiens RAB33A, member RAS oncogene family (RAB33A), mRNA  NM_004813 Homo sapiens PET112-like (yeast) (PET112L), mRNA	NM 004617	Homo sapiens transmembrane 4 superfamily member 4 (TM4SF4), mRNA
NM_004602 Homo sapiens staufen, RNA binding protein (Drosophila) (STAU), transcript variant T4, mRNA  NM_004653 Homo sapiens Smcy homolog, Y chromosome (mouse) (SMCY), mRNA  NM_004787 Homo sapiens slit homolog 2 (Drosophila) (SLIT2), mRNA  NM_004593 Homo sapiens splicing factor, arginine/serine-rich 10 (transformer 2 homolog, Drosophila) (SFRS10), mRNA  NM_004206 Homo sapiens vesicle trafficking protein (SEC22C), transcript variant 2, mRNA  NM_004657 Homo sapiens serum deprivation response (phosphatidylserine binding protein) (SDPR), mRNA  NM_004589 Homo sapiens SCO cytochrome oxidase deficient homolog 1 (yeast) (SCO1), nuclear gene encoding mitochondrial protein, mRNA  NM_004587 Homo sapiens ribosome binding protein 1 homolog 180kD (dog) (RRBP1), mRNA  NM_004164 Homo sapiens retinol binding protein 2, cellular (RBP2), mRNA  NM_004584 Homo sapiens RAD9 homolog (S. pombe) (RAD9), mRNA  NM_004794 Homo sapiens RAB33A, member RAS oncogene family (RAB33A), mRNA  NM_004813 Homo sapiens PET112-like (yeast) (PET112L), mRNA	NM 004607	Homo sapiens tubulin-specific chaperone a (TBCA), mRNA
NM_004653 Homo sapiens Smcy homolog, Y chromosome (mouse) (SMCY), mRNA NM_004787 Homo sapiens slit homolog 2 (Drosophila) (SLIT2), mRNA NM_004593 Homo sapiens splicing factor, arginine/serine-rich 10 (transformer 2 homolog, Drosophila) (SFRS10), mRNA NM_004206 Homo sapiens vesicle trafficking protein (SEC22C), transcript variant 2, mRNA NM_004657 Homo sapiens serum deprivation response (phosphatidylserine binding protein) (SDPR), mRNA NM_004589 Homo sapiens SCO cytochrome oxidase deficient homolog 1 (yeast) (SCO1), nuclear gene encoding mitochondrial protein, mRNA NM_004587 Homo sapiens ribosome binding protein 1 homolog 180kD (dog) (RRBP1), mRNA NM_004164 Homo sapiens retinol binding protein 2, cellular (RBP2), mRNA NM_004584 Homo sapiens RAD9 homolog (S. pombe) (RAD9), mRNA NM_004794 Homo sapiens RAB33A, member RAS oncogene family (RAB33A), mRNA NM_004813 Homo sapiens peroxisomal biogenesis factor 16 (PEX16), transcript variant 1, mRNA NM_004564 Homo sapiens PET112-like (yeast) (PET112L), mRNA	NM_004602	Homo sapiens staufen, RNA binding protein (Drosophila) (STAU), transcript
NM_004587 Homo sapiens slit homolog 2 (Drosophila) (SLIT2), mRNA  NM_004593 Homo sapiens splicing factor, arginine/serine-rich 10 (transformer 2 homolog, Drosophila) (SFRS10), mRNA  NM_004206 Homo sapiens vesicle trafficking protein (SEC22C), transcript variant 2, mRNA  NM_004657 Homo sapiens serum deprivation response (phosphatidylserine binding protein) (SDPR), mRNA  NM_004589 Homo sapiens SCO cytochrome oxidase deficient homolog 1 (yeast) (SCO1), nuclear gene encoding mitochondrial protein, mRNA  NM_004587 Homo sapiens ribosome binding protein 1 homolog 180kD (dog) (RRBP1), mRNA  NM_004164 Homo sapiens retinol binding protein 2, cellular (RBP2), mRNA  NM_004584 Homo sapiens RAD9 homolog (S. pombe) (RAD9), mRNA  NM_004794 Homo sapiens RAB33A, member RAS oncogene family (RAB33A), mRNA  NM_004813 Homo sapiens peroxisomal biogenesis factor 16 (PEX16), transcript variant 1, mRNA  NM_004564 Homo sapiens PET112-like (yeast) (PET112L), mRNA	NM 004653	
NM_004593 Homo sapiens splicing factor, arginine/serine-rich 10 (transformer 2 homolog, Drosophila) (SFRS10), mRNA  NM_004206 Homo sapiens vesicle trafficking protein (SEC22C), transcript variant 2, mRNA  NM_004657 Homo sapiens serum deprivation response (phosphatidylserine binding protein) (SDPR), mRNA  NM_004589 Homo sapiens SCO cytochrome oxidase deficient homolog 1 (yeast) (SC01), nuclear gene encoding mitochondrial protein, mRNA  NM_004587 Homo sapiens ribosome binding protein 1 homolog 180kD (dog) (RRBP1), mRNA  NM_004164 Homo sapiens retinol binding protein 2, cellular (RBP2), mRNA  NM_004584 Homo sapiens RAD9 homolog (S. pombe) (RAD9), mRNA  NM_004794 Homo sapiens RAB33A, member RAS oncogene family (RAB33A), mRNA  NM_004813 Homo sapiens PET112-like (yeast) (PET112L), mRNA		
NM_004206 Homo sapiens vesicle trafficking protein (SEC22C), transcript variant 2, mRNA  NM_004657 Homo sapiens serum deprivation response (phosphatidylserine binding protein) (SDPR), mRNA  NM_004589 Homo sapiens SCO cytochrome oxidase deficient homolog 1 (yeast) (SCO1), nuclear gene encoding mitochondrial protein, mRNA  NM_004587 Homo sapiens ribosome binding protein 1 homolog 180kD (dog) (RRBP1), mRNA  NM_004164 Homo sapiens retinol binding protein 2, cellular (RBP2), mRNA  NM_004584 Homo sapiens RAD9 homolog (S. pombe) (RAD9), mRNA  NM_004794 Homo sapiens RAB33A, member RAS oncogene family (RAB33A), mRNA  NM_004813 Homo sapiens peroxisomal biogenesis factor 16 (PEX16), transcript variant 1, mRNA  NM_004564 Homo sapiens PET112-like (yeast) (PET112L), mRNA	NM_004593	Homo sapiens splicing factor, arginine/serine-rich 10 (transformer 2 homolog,
NM_004589 Homo sapiens SCO cytochrome oxidase deficient homolog 1 (yeast) (SCO1), nuclear gene encoding mitochondrial protein, mRNA  NM_004587 Homo sapiens SCO cytochrome oxidase deficient homolog 1 (yeast) (SCO1), nuclear gene encoding mitochondrial protein, mRNA  NM_004587 Homo sapiens ribosome binding protein 1 homolog 180kD (dog) (RRBP1), mRNA  NM_004164 Homo sapiens retinol binding protein 2, cellular (RBP2), mRNA  NM_004584 Homo sapiens RAD9 homolog (S. pombe) (RAD9), mRNA  NM_004794 Homo sapiens RAB33A, member RAS oncogene family (RAB33A), mRNA  NM_004813 Homo sapiens peroxisomal biogenesis factor 16 (PEX16), transcript variant 1, mRNA  NM_004564 Homo sapiens PET112-like (yeast) (PET112L), mRNA	NM_004206	
NM_004589 Homo sapiens SCO cytochrome oxidase deficient homolog 1 (yeast) (SCO1), nuclear gene encoding mitochondrial protein, mRNA  NM_004587 Homo sapiens ribosome binding protein 1 homolog 180kD (dog) (RRBP1), mRNA  NM_004164 Homo sapiens retinol binding protein 2, cellular (RBP2), mRNA  NM_004584 Homo sapiens RAD9 homolog (S. pombe) (RAD9), mRNA  NM_004794 Homo sapiens RAB33A, member RAS oncogene family (RAB33A), mRNA  NM_004813 Homo sapiens peroxisomal biogenesis factor 16 (PEX16), transcript variant 1, mRNA  NM_004564 Homo sapiens PET112-like (yeast) (PET112L), mRNA	NM_004657	Homo sapiens serum deprivation response (phosphatidylserine binding protein)
NM_004587 Homo sapiens ribosome binding protein 1 homolog 180kD (dog) (RRBP1), mRNA  NM_004164 Homo sapiens retinol binding protein 2, cellular (RBP2), mRNA  NM_004584 Homo sapiens RAD9 homolog (S. pombe) (RAD9), mRNA  NM_004794 Homo sapiens RAB33A, member RAS oncogene family (RAB33A), mRNA  NM_004813 Homo sapiens peroxisomal biogenesis factor 16 (PEX16), transcript variant 1, mRNA  NM_004564 Homo sapiens PET112-like (yeast) (PET112L), mRNA	NM_004589	Homo sapiens SCO cytochrome oxidase deficient homolog 1 (yeast) (SCO1),
NM_004164Homo sapiens retinol binding protein 2, cellular (RBP2), mRNANM_004584Homo sapiens RAD9 homolog (S. pombe) (RAD9), mRNANM_004794Homo sapiens RAB33A, member RAS oncogene family (RAB33A), mRNANM_004813Homo sapiens peroxisomal biogenesis factor 16 (PEX16), transcript variant 1, mRNANM_004564Homo sapiens PET112-like (yeast) (PET112L), mRNA	NM_004587	Homo sapiens ribosome binding protein 1 homolog 180kD (dog) (RRBP1),
NM_004584 Homo sapiens RAD9 homolog (S. pombe) (RAD9), mRNA  NM_004794 Homo sapiens RAB33A, member RAS oncogene family (RAB33A), mRNA  NM_004813 Homo sapiens peroxisomal biogenesis factor 16 (PEX16), transcript variant 1, mRNA  NM_004564 Homo sapiens PET112-like (yeast) (PET112L), mRNA	NM 004164	
NM_004794 Homo sapiens RAB33A, member RAS oncogene family (RAB33A), mRNA NM_004813 Homo sapiens peroxisomal biogenesis factor 16 (PEX16), transcript variant 1, mRNA NM_004564 Homo sapiens PET112-like (yeast) (PET112L), mRNA		Homo sapiens RAD9 homolog (S. nombe) (PADO) DNA
NM_004813 Homo sapiens peroxisomal biogenesis factor 16 (PEX16), transcript variant 1, mRNA  NM_004564 Homo sapiens PET112-like (yeast) (PET112L), mRNA		
mRNA NM_004564 Homo sapiens PET112-like (yeast) (PET112L), mRNA		
		mRNA
NM_004643   Homo sapiens poly(A) binding protein, nuclear 1 (PABPN1), mRNA		
	NM_004643	Homo sapiens poly(A) binding protein, nuclear 1 (PABPN1), mRNA

NM_004561	Homo sapiens ovo-like 1(Drosophila) (OVOL1), mRNA
NM_004153	Homo sapiens origin recognition complex, subunit 1-like (yeast) (ORC1L),
_	mRNA
NM_004557	Homo sapiens Notch homolog 4 (Drosophila) (NOTCH4), mRNA
NM 004808	Homo sapiens N-myristoyltransferase 2 (NMT2), mRNA
NM 004210	Homo sapiens neuralized-like (Drosophila) (NEURL), mRNA
NM_004147	Homo sapiens developmentally regulated GTP binding protein 1 (DRG1),
	mRNA
NM_004851	Homo sapiens pronapsin A (NAP1), mRNA
NM_004533	Homo sapiens myosin binding protein C, fast type (MYBPC2), mRNA
NM_004529	Homo sapiens myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog,
	Drosophila); translocated to, 3 (MLLT3), mRNA
NM_004668	Homo sapiens maltase-glucoamylase (alpha-glucosidase) (MGAM), mRNA
NM_004526	Homo sapiens MCM2 minichromosome maintenance deficient 2, mitotin (S.
	cerevisiae) (MCM2), mRNA
NM_004829	Homo sapiens lymphocyte antigen 94 homolog, activating NK-receptor; NK-
	p46, (mouse) (LY94), mRNA
NM_004744	Homo sapiens lecithin retinol acyltransferase (phosphatidylcholineretinol O-
	acyltransferase) (LRAT), mRNA
NM_004524	Homo sapiens lethal giant larvae homolog 2 (Drosophila) (LLGL2), mRNA
NM_004140	Homo sapiens lethal giant larvae homolog 1 (Drosophila) (LLGL1), mRNA
NM_004922	Homo sapiens SEC24 related gene family, member C (S. cerevisiae) (SEC24C),
	mRNA
NM_004508	Homo sapiens isopentenyl-diphosphate delta isomerase (IDI1), mRNA
NM_004507	Homo sapiens HUS1 checkpoint homolog (S. pombe) (HUS1), mRNA
NM_004262	Homo sapiens airway trypsin-like protease (HAT), mRNA
NM_004752	Homo sapiens glial cells missing homolog b (Drosophila) (GCMB), mRNA
NM_004477	Homo sapiens FSHD region gene 1 (FRG1), mRNA
NM_004463	Homo sapiens faciogenital dysplasia (Aarskog-Scott syndrome) (FGD1), mRNA
NM_004106	Homo sapiens Fc fragment of IgE, high affinity I, receptor for; gamma polypeptide (FCER1G), mRNA
NM 004456	Homo sapiens enhancer of zeste homolog 2 (Drosophila) (EZH2), mRNA
NM 004100	Homo sapiens eyes absent homolog 4 (Drosophila) (EYA4), mRNA
NM 004450	Homo sapiens enhancer of rudimentary homolog (Drosophila) (ERH), mRNA
NM 004448	Homo sapiens v-erb-b2 erythroblastic leukemia viral oncogene homolog 2,
_	neuro/glioblastoma derived oncogene homolog (avian) (ERBB2), mRNA
NM 004445	Homo sapiens EphB6 (EPHB6), mRNA
NM 004436	Homo sapiens endosulfine alpha (ENSA), mRNA
NM 004432	Homo sapiens ELAV (embryonic lethal, abnormal vision, Drosophila)-like 2 (Hu
_	antigen B) (ELAVL2), mRNA
NM 004230	Homo sapiens endothelial differentiation, sphingolipid G-protein-coupled
_	receptor, 5 (EDG5), mRNA
NM 004421	Homo sapiens dishevelled, dsh homolog 1 (Drosophila) (DVL1), mRNA
NM 004399	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 11 (CHL1-like
_	helicase homològ, S. cerevisiae) (DDX11), transcript variant 2, mRNA
NM 004378	Homo sapiens cellular retinoic acid binding protein 1 (CRABP1), mRNA
NM 004898	Homo sapiens clock homolog (mouse) (CLOCK), mRNA
NM 004669	Homo sapiens chloride intracellular channel 3 (CLIC3), mRNA
NM 004066	Homo sapiens centrin, EF-hand protein, 1 (CETN1), mRNA
NM 004354	Homo sapiens cyclin G2 (CCNG2), mRNA
NM 004352	Homo sapiens cerebellin 1 precursor (CBLN1), mRNA
NM 004057	Homo sapiens calbindin 3, (vitamin D-dependent calcium binding protein)
14141 004021	1 Alonio sapions caronium 5, (vitamin 5 dependent calcium onium grotem)

	(CALB3), mRNA
NM_004338	Homo sapiens chromosome 18 open reading frame 1 (C18orf1), mRNA
NM_004725	Homo sapiens BUB3 budding uninhibited by benzimidazoles 3 homolog (yeast) (BUB3), mRNA
NM_004336	Homo sapiens BUB1 budding uninhibited by benzimidazoles 1 homolog (yeast) (BUB1), mRNA
NM_004331	Homo sapiens BCL2/adenovirus E1B 19kD interacting protein 3-like (BNIP3L), mRNA
NM 004328	Homo sapiens BCS1-like (yeast) (BCS1L), mRNA
NM 004045	Homo sapiens ATX1 antioxidant protein 1 homolog (yeast) (ATOX1), mRNA
NM_004849	Homo sapiens APG5 autophagy 5-like (S. cerevisiae) (APG5L), mRNA  Homo sapiens ash2 (absent, small, or homeotic)-like (Drosophila) (ASH2L),
NM_004674	mRNA
NM_004316	Homo sapiens achaete-scute complex-like 1 (Drosophila) (ASCL1), mRNA
NM_004707	Homo sapiens APG12 autophagy 12-like (S. cerevisiae) (APG12L), mRNA
NM_004641	Homo sapiens myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 10 (MLLT10), mRNA
NM 004301	Homo sapiens BAF53 (BAF53A), mRNA
NM 001129	Homo sapiens AE binding protein 1 (AEBP1), mRNA
NM 003656	Homo sapiens calcium/calmodulin-dependent protein kinase I (CAMK1), mRNA
NM 000239	Homo sapiens lysozyme (renal amyloidosis) (LYZ), mRNA
NM 000456	Homo sapiens sulfite oxidase (SUOX), nuclear gene encoding mitochondrial
11112_000 100	protein, mRNA
NM_000435	Homo sapiens Notch homolog 3 (Drosophila) (NOTCH3), mRNA
NM 000251	Homo sapiens mutS homolog 2, colon cancer, nonpolyposis type 1 (E. coli)
	(MSH2), mRNA
NM_000249	Homo sapiens mutL homolog 1, colon cancer, nonpolyposis type 2 (E. coli) (MLH1), mRNA
NM 000210	Homo sapiens integrin, alpha 6 (ITGA6), mRNA
NM 001537	Homo sapiens heat shock factor binding protein 1 (HSBP1), mRNA
NM 001499	Homo sapiens GLE1 RNA export mediator-like (yeast) (GLE1L), mRNA
NM 001458	Homo sapiens filamin C, gamma (actin binding protein 280) (FLNC), mRNA
NM_001444	Homo sapiens fatty acid binding protein 5 (psoriasis-associated) (FABP5), mRNA
NM 001432	Homo sapiens epiregulin (EREG), mRNA
NM_001388	Homo sapiens developmentally regulated GTP binding protein 2 (DRG2), mRNA
NM_001340	Homo sapiens cylicin, basic protein of sperm head cytoskeleton 2 (CYLC2), mRNA
NM_001326	Homo sapiens cleavage stimulation factor, 3' pre-RNA, subunit 3, 77kD (CSTF3), mRNA
NM_001325	Homo sapiens cleavage stimulation factor, 3' pre-RNA, subunit 2, 64kD (CSTF2), mRNA
NM_001324	Homo sapiens cleavage stimulation factor, 3' pre-RNA, subunit 1, 50kD (CSTF1), mRNA
NM_001255	Homo sapiens CDC20 cell division cycle 20 homolog (S. cerevisiae) (CDC20), mRNA
NM 001122	Homo sapiens adipose differentiation-related protein (ADFP), mRNA
NM 003413	Homo sapiens Zic family member 3 heterotaxy 1 (odd-paired homolog,
	Drosophila) (ZIC3), mRNA
NM_003412	Homo sapiens Zic family member 1 (odd-paired homolog, Drosophila) (ZIC1), mRNA
L———————	1

NM_003408	Homo sapiens zinc finger protein 37 homolog (mouse) (ZFP37), mRNA
NM_003409	Homo sapiens zinc finger protein 161 homolog (mouse) (ZFP161), mRNA
NM_003680	Homo sapiens tyrosyl-tRNA synthetase (YARS), mRNA
NM_003390	Homo sapiens WEE1+ homolog (S. pombe) (WEE1), mRNA
NM_003565	Homo sapiens unc-51-like kinase 1 (C. elegans) (ULK1), mRNA
NM_003345	Homo sapiens ubiquitin-conjugating enzyme E2I (UBC9 homolog, yeast) (UBE2I), mRNA
NM_003344	Homo sapiens ubiquitin-conjugating enzyme E2H (UBC8 homolog, yeast) (UBE2H), mRNA
NM_003343	Homo sapiens ubiquitin-conjugating enzyme E2G 2 (UBC7 homolog, yeast) (UBE2G2), mRNA
NM_003340	Homo sapiens ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homolog, yeast) (UBE2D3), mRNA
NM_003338	Homo sapiens ubiquitin-conjugating enzyme E2D 1 (UBC4/5 homolog, yeast) (UBE2D1), mRNA
NM_003968	Homo sapiens ubiquitin-activating enzyme E1C (UBA3 homolog, yeast) (UBE1C), mRNA
NM_003320	Homo sapiens tubby homolog (mouse) (TUB), mRNA
NM_003278	Homo sapiens tetranectin (plasminogen binding protein) (TNA), mRNA
NM_003260	Homo sapiens transducin-like enhancer of split 2 (E(sp1) homolog, Drosophila) (TLE2), mRNA
NM_003920	Homo sapiens timeless homolog (Drosophila) (TIMELESS), mRNA
NM_003251	Homo sapiens thyroid hormone responsive (SPOT14 homolog, rat) (THRSP), mRNA
NM_003250	Homo sapiens thyroid hormone receptor, alpha (erythroblastic leukemia viral (verb-a) oncogene homolog, avian) (THRA), mRNA
NM_003223	Homo sapiens transcription factor AP-4 (activating enhancer binding protein 4) (TFAP4), mRNA
NM_003222	Homo sapiens transcription factor AP-2 gamma (activating enhancer binding protein 2 gamma) (TFAP2C), mRNA
NM_003221	Homo sapiens transcription factor AP-2 beta (activating enhancer binding protein 2 beta) (TFAP2B), mRNA
NM_003220	Homo sapiens transcription factor AP-2 alpha (activating enhancer binding protein 2 alpha) (TFAP2A), mRNA
NM_000458	Homo sapiens transcription factor 2, hepatic; LF-B3; variant hepatic nuclear factor (TCF2), transcript variant a, mRNA
NM_003181	Homo sapiens T, brachyury homolog (mouse) (T), mRNA
NM_003173	Homo sapiens suppressor of variegation 3-9 homolog 1 (Drosophila) (SUV39H1), mRNA
NM_003171	Homo sapiens suppressor of var1, 3-like 1 (S. cerevisiae) (SUPV3L1), mRNA
NM_003169	Homo sapiens suppressor of Ty 5 homolog (S. cerevisiae) (SUPT5H), mRNA
NM_003168	Homo sapiens suppressor of Ty 4 homolog 1 (S. cerevisiae) (SUPT4H1), mRNA
NM_003599	Homo sapiens suppressor of Ty 3 homolog (S. cerevisiae) (SUPT3H), mRNA
NM_003162	Homo sapiens striatin, calmodulin binding protein (STRN), mRNA
NM_003134	Homo sapiens signal recognition particle 14kD (homologous Alu RNA binding protein) (SRP14), mRNA
NM_003088	Homo sapiens singed-like (fascin homolog, sea urchin) (Drosophila) (SNL), mRNA
NM_003061	Homo sapiens slit homolog 1 (Drosophila) (SLIT1), mRNA
NM_003036	Homo sapiens v-ski sarcoma viral oncogene homolog (avian) (SKI), mRNA
NM_003031	Homo sapiens seven in absentia homolog 1 (Drosophila) (SIAH1), mRNA
NM_000193	Homo sapiens sonic hedgehog homolog (Drosophila) (SIAH), mRNA
	- B Moniore (Drosopinia) (SIIII), IIININA

NM_003003	Homo sapiens SEC14-like 1 (S. cerevisiae) (SEC14L1), mRNA
NM_002983	Homo sapiens small inducible cytokine A3 (SCYA3), mRNA
NM_002982	Homo sapiens small inducible cytokine A2 (monocyte chemotactic protein 1) (SCYA2), mRNA
NM_002981	Homo sapiens small inducible cytokine A1, I-309 (SCYA1), mRNA
NM_003864	Homo sapiens sin3-associated polypeptide, 30kD (SAP30), mRNA
NM_002962	Homo sapiens S100 calcium binding protein A5 (S100A5), mRNA
NM_002960	Homo sapiens S100 calcium binding protein A3 (S100A3), mRNA
NM_002966	Homo sapiens S100 calcium binding protein A10 (annexin II ligand, calpactin I, light polypeptide (p11)) (S100A10), mRNA
NM_003707	Homo sapiens RuvB-like 1 (E. coli) (RUVBL1), mRNA
NM_002944	Homo sapiens v-ros UR2 sarcoma virus oncogene homolog 1 (avian) (ROS1), mRNA
NM_002941	Homo sapiens roundabout, axon guidance receptor, homolog 1 (Drosophila) (ROBO1), mRNA
NM_000326	Homo sapiens retinaldehyde binding protein 1 (RLBP1), mRNA
NM_002930	Homo sapiens Ric-like, expressed in neurons (Drosophila) (RIN), mRNA
NM_003961	Homo sapiens rhomboid, veinlet-like 1 (Drosophila) (RHBDL), mRNA
NM_002912	Homo sapiens REV3-like, catalytic subunit of DNA polymerase zeta (yeast) (REV3L), mRNA
NM_002900	Homo sapiens retinol binding protein 3, interstitial (RBP3), mRNA
NM_002894	Homo sapiens retinoblastoma binding protein 8 (RBBP8), mRNA
NM_002888	Homo sapiens retinoic acid receptor responder (tazarotene induced) 1 (RARRES1), mRNA
NM_002879	Homo sapiens RAD52 homolog (S. cerevisiae) (RAD52) mRNA
NM_002878	Homo sapiens RAD51-like 3 (S. cerevisiae) (RAD51L3), mRNA
NM_002875	Homo sapiens RAD51 homolog (RecA homolog, E. coli) (S. cerevisiae) (RAD51), mRNA
NM_002874	Homo sapiens RAD23 homolog B (S. cerevisiae) (RAD23B), mRNA
NM_002853	Homo sapiens RAD1 homolog (S. pombe) (RAD1), mRNA
NM_002873	Homo sapiens RAD17 homolog (S. pombe) (RAD17), mRNA
NM_000264	Homo sapiens patched homolog (Drosophila) (PTCH), mRNA
NM_003738	Homo sapiens patched homolog 2 (Drosophila) (PTCH2), mRNA
NM_002616	Homo sapiens period homolog 1 (Drosophila) (PER1), mRNA
NM_002600	Homo sapiens phosphodiesterase 4B, cAMP-specific (phosphodiesterase E4
	dunce homolog, Drosophila) (PDE4B), mRNA
NM_002568	Homo sapiens poly(A) binding protein, cytoplasmic 1 (PABPC1), mRNA
NM_003932	Homo sapiens suppression of tumorigenicity 13 (colon carcinoma) (Hsp70 interacting protein) (ST13), mRNA
NM_003715	Homo sapiens vesicle docking protein p115 (P115), mRNA
NM_002553	Homo sapiens origin recognition complex, subunit 5-like (yeast) (ORC5L), mRNA
NM_002552	Homo sapiens origin recognition complex, subunit 4-like (yeast) (ORC4L), mRNA
NM_003634	Homo sapiens nipsnap homolog 1 (C. elegans) (NIPSNAP1), mRNA
NM_002499	Homo sapiens neogenin homolog 1 (chicken) (NEO1), mRNA
NM_002484	Homo sapiens nucleotide binding protein 1 (MinD homolog, E. coli) (NUBP1), mRNA
NM_003827	Homo sapiens N-ethylmaleimide-sensitive factor attachment protein, alpha (NAPA), mRNA
NM_002466	Homo sapiens v-myb myeloblastosis viral oncogene homolog (avian)-like 2 (MYBL2), mRNA

NM 002448	Homo sapiens msh homeo box homolog 1 (Drosophila) (MSX1), mRNA
NM 003576	Homo sapiens serine/threonine kinase 24 (STE20 homolog, yeast) (STK24),
1111_003370	mRNA
NM 002442	Homo sapiens musashi homolog 1 (Drosophila) (MSI1), mRNA
NM 002441	Homo sapiens mutS homolog 5 (E. coli) (MSH5), mRNA
NM 002440	Homo sapiens mutS homolog 4 (E. coli) (MSH4), mRNA
NM 002439	Homo sapiens mutS homolog 3 (E. coli) (MSH3), mRNA
NM 002405	Homo sapiens manic fringe homolog (Drosophila) (MFNG), mRNA
NM 002402	Homo sapiens mesoderm specific transcript homolog (mouse) (MEST), mRNA
NM_002398	Homo sapiens Meis1, myeloid ecotropic viral integration site 1 homolog (mouse)
_	(MEIS1), mRNA
NM_002393	Homo sapiens Mdm4, transformed 3T3 cell double minute 4, p53 binding protein
	(mouse) (MDM4), mRNA
NM_002392	Homo sapiens Mdm2, transformed 3T3 cell double minute 2, p53 binding protein
	(mouse) (MDM2), transcript variant MDM2, mRNA
NM_003906	Homo sapiens MCM3 minichromosome maintenance deficient 3 (S. cerevisiae)
	associated protein (MCM3AP), mRNA
NM_002360	Homo sapiens v-maf musculoaponeurotic fibrosarcoma oncogene homolog K
	(avian) (MAFK), mRNA
NM_002359	Homo sapiens v-maf musculoaponeurotic fibrosarcoma oncogene homolog G
) T ( 000 - 000	(avian) (MAFG), mRNA
NM_003550	Homo sapiens MAD1 mitotic arrest deficient-like 1 (yeast) (MAD1L1), mRNA
NM_003937	Homo sapiens kynureninase (L-kynurenine hydrolase) (KYNU), mRNA
NM_002269	Homo sapiens karyopherin alpha 5 (importin alpha 6) (KPNA5), mRNA
NM_003772	Homo sapiens jerky homolog-like (mouse) (JRKL), mRNA
NM_002202	Homo sapiens ISL1 transcription factor, LIM/homeodomain, (islet-1) (ISL1),
) D ( 000 (0 )	mRNA
NM_003604	Homo sapiens insulin receptor substrate 4 (IRS4), mRNA
NM_001570	Homo sapiens interleukin-1 receptor-associated kinase 2 (IRAK2), mRNA
NM_003866	Homo sapiens inositol polyphosphate-4-phosphatase, type II, 105kD (INPP4B), mRNA
NM_001536	Homo sapiens HMT1 hnRNP methyltransferase-like 2 (S. cerevisiae)
	(HRMT1L2), mRNA
NM_001535	Homo sapiens HMT1 hnRNP methyltransferase-like 1 (S. cerevisiae) (HRMT1L1), mRNA
NM_003806	Homò sapiens harakiri, BCL2 interacting protein (contains only BH3 domain)
	(HRK), mRNA
NM_002152	Homo sapiens histidine rich calcium binding protein (HRC), mRNA
NM_002114	Homo sapiens human immunodeficiency virus type I enhancer binding protein 1 (HIVEP1), mRNA
NM 003710	Homo sapiens serine protease inhibitor, Kunitz type 1 (SPINT1), mRNA
NM_000179	Homo sapiens mutS homolog 6 (E. coli) (MSH6), mRNA
NM 000839	Homo sapiens glutamate receptor, metabotropic 2 (GRM2), mRNA
NM 002077	Homo sapiens golgi autoantigen, golgin subfamily a, 1 (GOLGA1), mRNA
NM 003878	Homo sapiens gamma-glutamyl hydrolase (conjugase, folylpolygammaglutamyl
	hydrolase) (GGH), mRNA
NM_001488	Homo sapiens transcriptional adaptor 2 (ADA2 homolog, yeast)-like (TADA2L), mRNA
NM 001487	Homo sapiens GCN5 general control of amino-acid synthesis 5-like 1 (yeast)
- 12.2_001.107	(GCN5L1), mRNA
NM_003643	Homo sapiens glial cells missing homolog a (Drosophila) (GCMA), mRNA
NM 002052	Homo sapiens GATA binding protein 4 (GATA4), mRNA
	1 And the september Street billing protein 7 (OATA4), Indian

ND 6 000051	
NM_002051	Homo sapiens GATA binding protein 3 (GATA3), mRNA
NM_002050	Homo sapiens GATA binding protein 2 (GATA2), mRNA
NM_002049	Homo sapiens GATA binding protein 1 (globin transcription factor 1) (GATA1), mRNA
NM_002040	Homo sapiens GA binding protein transcription factor, alpha subunit (60kD) (GABPA), mRNA
NM 002039	Homo sapiens GRB2-associated binding protein 1 (GAB1), mRNA
NM_003508	Homo sapiens frizzled homolog 9 (Drosophila) (FZD9), mRNA
NM 003507	Homo sapiens frizzled homolog 7 (Drosophila) (FZD7), mRNA
NM 003506	Homo sapiens frizzled homolog 6 (Drosophila) (FZD6), mRNA
NM 003468	Homo sapiens frizzled homolog 5 (Drosophila) (FZD5), mRNA
NM 003505	Homo sapiens frizzled homolog 1 (Drosophila) (FZD1), mRNA
NM 001465	Homo sapiens FYN binding protein (FYB-120/130) (FYB), mRNA
NM 002031	Homo sapiens fyn-related kinase (FRK), mRNA
NM 003717	Homo sapiens neuropeptide FF-amide peptide precursor (NPFF), mRNA
NM 001457	Homo sapiens filamin B, beta (actin binding protein 278) (FLNB), mRNA
NM 001456	Homo sapiens filamin A, alpha (actin binding protein 280) (FLNA), mRNA
NM 002018	Homo sapiens flightless I homolog (Drosophila) (FLII), mRNA
NM 001991	Homo sapiens enhancer of zeste homolog 1 (Drosophila) (EZH1), mRNA
NM 001990	Homo sapiens eyes absent homolog 3 (Drosophila) (EYA3), mRNA
NM 000503	Homo sapiens eyes absent homolog 1 (Drosophila) (EYA1), mRNA
NM 001989	Homo sapiens eve, even-skipped homeo box homolog 1 (Drosophila) (EVX1),
_	mRNA
NM_001982	Homo sapiens v-erb-b2 erythroblastic leukemia viral oncogene homolog 3
	(avian) (ERBB3), mRNA
NM_003584	Homo sapiens dual specificity phosphatase 11 (RNA/RNP complex 1-
	interacting) (DUSP11), mRNA
NM_003859	Homo sapiens dolichyl-phosphate mannosyltransferase polypeptide 1, catalytic
	subunit (DPM1), mRNA
NM_001928	Homo sapiens D component of complement (adipsin) (DF), mRNA
NM_003649	Homo sapiens D-aspartate oxidase (DDO), transcript variant 1, mRNA
NM_001343	Homo sapiens disabled homolog 2, mitogen-responsive phosphoprotein
37.5 001010	(Drosophila) (DAB2), mRNA
NM_001913	Homo sapiens cut-like 1, CCAAT displacement protein (Drosophila) (CUTL1), mRNA
NM_001316	Homo sapiens CSE1 chromosome segregation 1-like (yeast) (CSE1L), mRNA
NM_003652	Homo sapiens carboxypeptidase Z (CPZ), mRNA
NM_003909	Homo sapiens copine III (CPNE3), mRNA
NM_003915	Homo sapiens copine I (CPNE1), mRNA
NM_001308	Homo sapiens carboxypeptidase N, polypeptide 1, 50kD (CPN1), mRNA
NM_001841	Homo sapiens cannabinoid receptor 2 (macrophage) (CNR2), mRNA
NM_001280	Homo sapiens cold inducible RNA binding protein (CIRBP), mRNA
NM_001274	Homo sapiens CHK1 checkpoint homolog (S. pombe) (CHEK1), mRNA
NM_001806	Homo sapiens CCAAT/enhancer binding protein (C/EBP), gamma (CEBPG), mRNA
NM_003655	Homo sapiens chromobox homolog 4 (Pc class homolog, Drosophila) (CBX4),
	mRNA
NM_001749	Homo sapiens calpain, small subunit 1 (CAPNS1), mRNA
NM_000716	Homo sapiens complement component 4 binding protein, beta (C4BPB), mRNA
NM_000715	Homo sapiens complement component 4 binding protein, alpha (C4BPA), mRNA
NM_001726	Homo sapiens bromodomain, testis-specific (BRDT), mRNA

NM_001205	Homo sapiens BCL2/adenovirus E1B 19kD interacting protein 1 (BNIP1), transcript variant BNIP1, mRNA
NM 001714	Homo sapiens Bicaudal D homolog 1 (Drosophila) (BICD1), mRNA
NM 003766	Homo sapiens beclin 1 (coiled-coil, myosin-like BCL2 interacting protein)
1414_003700	(BECN1), mRNA
NM 003567	Homo sapiens breast cancer anti-estrogen resistance 3 (BCAR3), mRNA
NM 001189	Homo sapiens bagpipe homeobox homolog 1 (Drosophila) (BAPX1), mRNA
NM 001698	Homo sapiens AU RNA binding protein/enoyl-Coenzyme A hydratase (AUH),
_	nuclear gene encoding mitochondrial protein, mRNA
NM_001672	Homo sapiens agouti signaling protein, nonagouti homolog (mouse) (ASIP), mRNA
NM 001638	Homo sapiens apolipoprotein F (APOF), mRNA
NM 003977	Homo sapiens aryl hydrocarbon receptor interacting protein (AIP), mRNA
NM_001138	Homo sapiens agouti related protein homolog (mouse) (AGRP), transcript variant 1, mRNA
NM_058246	Homo sapiens DnaJ (Hsp40) homolog, subfamily B, member 6 (DNAJB6), mRNA
NM 025225	Homo sapiens hypothetical protein dJ796I17.1 (DJ796I17.1), mRNA
NM_058165	Homo sapiens diacylglycerol acyltransferase 2-like (DGAT2-like), mRNA
NM 001861	Homo sapiens cytochrome c oxidase subunit IV isoform 1 (COX4II), nuclear
	gene encoding mitochondrial protein, mRNA
NM 014491	Homo sapiens forkhead box P2 (FOXP2), mRNA
NM 054110	Homo sapiens UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-
<u>-</u>	acetylgalactosaminyltransferase 7 (GALNT7), mRNA
NM_006726	Homo sapiens vesicle trafficking, beach and anchor containing (LRBA), mRNA
NM_020663	Homo sapiens TC10-like Rho GTPase (TCL), mRNA
NM_020919	Homo sapiens amyotrophic lateral sclerosis 2 (juvenile) (ALS2), mRNA
NM_052852	Homo sapiens hypothetical zinc finger protein MGC2396 (MGC2396), mRNA
NM_053043	Homo sapiens hypothetical protein MGC20460 (MGC20460), mRNA
NM_053017	Homo sapiens ADP-ribosyltransferase 5 (ART5), mRNA
NM_052999	Homo sapiens chemokine-like factor-like protein CKLFH1 (CKLFH1), mRNA
NM_052881	Homo sapiens hypothetical protein dJ734P14.5 (novel C2H2 type zinc finger protein) (MGC20504), mRNA
NM 052968	Homo sapiens apolipoprotein A-V (APOA5), mRNA
NM 052960	Homo sapiens retinoid binding protein 7 (RBP7), mRNA
NM 052959	Homo sapiens pannexin 3 (PANX3), mRNA
NM 052948	Homo sapiens sorting nexin 26 (SNX26), mRNA
NM 052947	Homo sapiens heart alpha-kinase (HAK), mRNA
NM 052946	Homo sapiens hypothetical protein MGC20702 (MGC20702), mRNA
NM 052943	Homo sapiens hypothetical protein MGC16491 (MGC16491), mRNA
NM 052941	Homo sapiens guanylate binding protein 4 (GBP4), mRNA
NM 052935	Homo sapiens hypothetical protein MGC20781 (MGC20781), mRNA
NM 052890	Homo sapiens peptidoglycan recognition protein L precursor (PGLYRP), mRNA
NM_052885	Homo sapiens solute carrier family 2 (facilitated glucose transporter), member 13
	(SLC2A13), mRNA
NM_052884	Homo sapiens sialic acid binding Ig-like lectin 11 (SIGLEC11), mRNA
NM_052877	Homo sapiens similar to hypothetical protein MNCb-2386 (MGC17544), mRNA
NM_052876	Homo sapiens transcriptional repressor NAC1 (NAC1), mRNA
NM_052873	Homo sapiens MGC16028 similar to RIKEN cDNA 1700019E19 gene (MGC16028), mRNA
NM 052871	Homo sapiens hypothetical protein MGC4677 (MGC4677), mRNA
NM 052870	Homo sapiens sorting nexin 18 (SNX18), mRNA

NM_052859	Homo sapiens putative endoplasmic reticulum multispan transmembrane protein
NIM 052050	(RFT1), mRNA
NM_052858 NM_052855	Homo sapiens similar to RIKEN cDNA 1810006A16 gene (LOC91862), mRNA
	Homo sapiens hypothetical protein MGC15396 (MGC15396), mRNA
NM_052854	Homo sapiens old astrocyte specifically induced substance (OASIS), mRNA
NM_052844	Homo sapiens hypothetical protein MGC20486 (MGC20486), mRNA
NM_052839	Homo sapiens pannexin 2 (PANX2), mRNA
NM_033551 NM_033549	Homo sapiens hypothetical protein MGC19556 (MGC19556), mRNA
NM 033546	Homo sapiens hypothetical gene MGC1127 (MGC1127), mRNA
NM 033544	Homo sapiens myosin regulatory light chain (MLC-B), mRNA
NM_033344	Homo sapiens similar to cyclin-E binding protein 1 (H. sapiens) (MGC14386), mRNA
NM 033515	Homo sapiens MacGAP protein (MacGAP), mRNA
NM 033519	Homo sapiens olfactory receptor sdolf (sdolf), mRNA
NM 033516	Homo sapiens protein kinase NYD-SP25 (NYD-SP25), mRNA
NM 032231	Homo sapiens hypothetical protein FLJ22875 (FLJ22875), mRNA
NM 018437	Homo sapiens hypothetical protein EDAG-1 (EDAG-1), mRNA
NM 033378	Homo sapiens chorionic gonadotropin, beta polypeptide 2 (CGB2), mRNA
NM_033377	Homo sapiens chorionic gonadotropin, beta polypeptide 2 (CGB2), mRNA
NM 033448	Homo sapiens keratin 6 irs (KRT6IRS), mRNA
NM 033424	Homo sapiens similar to MYOSIN HEAVY CHAIN, CARDIAC MUSCLE
	ALPHA ISOFORM (MYHC-ALPHA) (M. musculus) (LOC92771), mRNA
NM 033445	Homo sapiens similar to H2A histone family, member A (H. sapiens)
_	(MGC3165), mRNA
NM 033439	Homo sapiens DVS27-related protein (DVS27), mRNA
NM 033440	Homo sapiens elastase 2A (ELA2A), mRNA
NM 033438	Homo sapiens CD84-H1 precursor (CD84-H1), mRNA
NM 033423	Homo sapiens similar to granzyme B (granzyme 2, cytotoxic T-lymphocyte-
-	associated serine esterase 1) (H. sapiens) (CTLA1), mRNA
NM_033411	Homo sapiens hypothetical protein MGC13523 (MGC13523), mRNA
NM_033416	Homo sapiens similar to HYPOTHETICAL 34.0 KDA PROTEIN ZK795.3 IN
	CHROMOSOME IV (MGC19606), mRNA
NM_033413	Homo sapiens hypothetical gene MGC16309 (MGC16309), mRNA
NM_033410	Homo sapiens hypothetical protein MGC13138 (MGC13138), mRNA
NM_033419	Homo sapiens hypothetical gene MGC9753 (MGC9753), mRNA
NM_014083	Homo sapiens PRO0767 protein (PRO0767), mRNA
NM_033043	Homo sapiens chorionic gonadotropin, beta polypeptide 5 (CGB5), mRNA
NM_031451	Homo sapiens hypothetical protein MGC4766 similar to testis specific protein
	TES101RP (MGC4766), mRNA
NM_033183	Homo sapiens chorionic gonadotropin, beta polypeptide 8 (CGB8), mRNA
NM_020443	Homo sapiens hypothetical protein MGC14961 (MGC14961), mRNA
NM_033343	Homo sapiens LIM homeobox protein 4 (LHX4), mRNA
NM_033318	Homo sapiens hypothetical gene supported by AL449243 (LOC91689), mRNA
NM_033328	Homo sapiens capping protein alpha 3 (CAPPA3), mRNA
NM_033315	Homo sapiens ras-like protein VTS58635 (VTS58635), mRNA
NM_033309	Homo sapiens hypothetical protein MGC4655 (MGC4655), mRNA
NM_033296	Homo sapiens T-cell activation protein (PGR1), mRNA
NM_033297	Homo sapiens leucine-rich-repeat protein (RNO2), mRNA
NM_033280	Homo sapiens similar to signal peptidase complex (18kD) (LOC90701), mRNA
NM_033196	Homo sapiens similar to ZINC FINGER PROTEIN 85 (ZINC FINGER
370.6.00	PROTEIN HPF4) (HTF1) (H. sapiens) (LOC91120), mRNA
NM_033272	Homo sapiens potassium channel subunit HERG-3 (HERG-3), mRNA

NM 033261 Homo sapiens diphosphate dimethylallyl diphosphate isomerase 2 (IDI2), mRN NM 033254 Homo sapiens brother of CDO (BOC), mRNA  NM 033204 Homo sapiens hypothetical gene DKFZp570I0164 (DKFZp570I0164), mRNA NM 033259 Homo sapiens CaM-KII inhibitory protein (CAM-KIIN), mRNA NM 032597 Homo sapiens testes development-related NYD-SP21 (NYD-SP21), mRNA NM 033212 Homo sapiens hypothetical gene supported by BC004307; BC008285 (MGC10992), mRNA  NM 033208 Homo sapiens similar to jerky (mouse) homolog-like (LOC91151), mRNA NM 033195 Homo sapiens lactate dehydrogenase A -like (LDHL), mRNA NM 015643 Homo sapiens DKFZP434F122 protein (DKFZP434F122), mRNA NM 032604 Homo sapiens lung alpha/beta hydrolase 1 (LABH1), mRNA NM 032133 Homo sapiens hypothetical protein DKFZp434N1415 (DKFZP434N1415), mRNA NM 030803 Homo sapiens hypothetical protein FLJ10035 (FLJ10035), mRNA NM 024062 Homo sapiens hypothetical protein MGC5338 (MGC5338), mRNA
NM_033204 Homo sapiens hypothetical gene DKFZp570I0164 (DKFZp570I0164), mRNA NM_033259 Homo sapiens CaM-KII inhibitory protein (CAM-KIIN), mRNA NM_032597 Homo sapiens testes development-related NYD-SP21 (NYD-SP21), mRNA NM_033212 Homo sapiens hypothetical gene supported by BC004307; BC008285 (MGC10992), mRNA NM_033208 Homo sapiens similar to jerky (mouse) homolog-like (LOC91151), mRNA NM_033195 Homo sapiens lactate dehydrogenase A -like (LDHL), mRNA NM_015643 Homo sapiens DKFZP434F122 protein (DKFZP434F122), mRNA NM_032604 Homo sapiens lung alpha/beta hydrolase 1 (LABH1), mRNA NM_032133 Homo sapiens hypothetical protein DKFZp434N1415 (DKFZP434N1415), mRNA NM_030803 Homo sapiens hypothetical protein FLJ10035 (FLJ10035), mRNA
NM 033259 Homo sapiens CaM-KII inhibitory protein (CAM-KIIN), mRNA  NM 032597 Homo sapiens testes development-related NYD-SP21 (NYD-SP21), mRNA  NM 033212 Homo sapiens hypothetical gene supported by BC004307; BC008285  (MGC10992), mRNA  NM 033208 Homo sapiens similar to jerky (mouse) homolog-like (LOC91151), mRNA  NM 033195 Homo sapiens lactate dehydrogenase A -like (LDHL), mRNA  NM 015643 Homo sapiens DKFZP434F122 protein (DKFZP434F122), mRNA  NM 032604 Homo sapiens lung alpha/beta hydrolase 1 (LABH1), mRNA  NM 032133 Homo sapiens hypothetical protein DKFZP434N1415 (DKFZP434N1415),  mRNA  NM 030803 Homo sapiens hypothetical protein FLJ10035 (FLJ10035), mRNA
NM_032597 Homo sapiens testes development-related NYD-SP21 (NYD-SP21), mRNA  NM_033212 Homo sapiens hypothetical gene supported by BC004307; BC008285 (MGC10992), mRNA  NM_033208 Homo sapiens similar to jerky (mouse) homolog-like (LOC91151), mRNA  NM_033195 Homo sapiens lactate dehydrogenase A -like (LDHL), mRNA  NM_015643 Homo sapiens DKFZP434F122 protein (DKFZP434F122), mRNA  NM_032604 Homo sapiens lung alpha/beta hydrolase 1 (LABH1), mRNA  NM_032133 Homo sapiens hypothetical protein DKFZp434N1415 (DKFZP434N1415),  mRNA  NM_030803 Homo sapiens hypothetical protein FLJ10035 (FLJ10035), mRNA
NM_033212 Homo sapiens hypothetical gene supported by BC004307; BC008285 (MGC10992), mRNA  NM_033208 Homo sapiens similar to jerky (mouse) homolog-like (LOC91151), mRNA  NM_033195 Homo sapiens lactate dehydrogenase A -like (LDHL), mRNA  NM_015643 Homo sapiens DKFZP434F122 protein (DKFZP434F122), mRNA  NM_032604 Homo sapiens lung alpha/beta hydrolase 1 (LABH1), mRNA  NM_032133 Homo sapiens hypothetical protein DKFZP434N1415 (DKFZP434N1415), mRNA  NM_030803 Homo sapiens hypothetical protein FLJ10035 (FLJ10035), mRNA
(MGC10992), mRNA  NM 033208 Homo sapiens similar to jerky (mouse) homolog-like (LOC91151), mRNA  NM 033195 Homo sapiens lactate dehydrogenase A -like (LDHL), mRNA  NM 015643 Homo sapiens DKFZP434F122 protein (DKFZP434F122), mRNA  NM 032604 Homo sapiens lung alpha/beta hydrolase 1 (LABH1), mRNA  NM 032133 Homo sapiens hypothetical protein DKFZp434N1415 (DKFZP434N1415), mRNA  NM 030803 Homo sapiens hypothetical protein FLJ10035 (FLJ10035), mRNA
NM_033208 Homo sapiens similar to jerky (mouse) homolog-like (LOC91151), mRNA  NM_033195 Homo sapiens lactate dehydrogenase A -like (LDHL), mRNA  NM_015643 Homo sapiens DKFZP434F122 protein (DKFZP434F122), mRNA  NM_032604 Homo sapiens lung alpha/beta hydrolase 1 (LABH1), mRNA  NM_032133 Homo sapiens hypothetical protein DKFZp434N1415 (DKFZP434N1415), mRNA  NM_030803 Homo sapiens hypothetical protein FLJ10035 (FLJ10035), mRNA
NM_033195 Homo sapiens lactate dehydrogenase A -like (LDHL), mRNA  NM_015643 Homo sapiens DKFZP434F122 protein (DKFZP434F122), mRNA  NM_032604 Homo sapiens lung alpha/beta hydrolase 1 (LABH1), mRNA  NM_032133 Homo sapiens hypothetical protein DKFZp434N1415 (DKFZP434N1415),  mRNA  NM_030803 Homo sapiens hypothetical protein FLJ10035 (FLJ10035), mRNA
NM_015643 Homo sapiens DKFZP434F122 protein (DKFZP434F122), mRNA  NM_032604 Homo sapiens lung alpha/beta hydrolase 1 (LABH1), mRNA  NM_032133 Homo sapiens hypothetical protein DKFZp434N1415 (DKFZP434N1415),  mRNA  NM_030803 Homo sapiens hypothetical protein FLJ10035 (FLJ10035), mRNA
NM_032604 Homo sapiens lung alpha/beta hydrolase 1 (LABH1), mRNA  NM_032133 Homo sapiens hypothetical protein DKFZp434N1415 (DKFZP434N1415),  mRNA  NM_030803 Homo sapiens hypothetical protein FLJ10035 (FLJ10035), mRNA
NM_032133 Homo sapiens hypothetical protein DKFZp434N1415 (DKFZP434N1415), mRNA  NM_030803 Homo sapiens hypothetical protein FLJ10035 (FLJ10035), mRNA
mRNA NM_030803 Homo sapiens hypothetical protein FLJ10035 (FLJ10035), mRNA
NM_030803 Homo sapiens hypothetical protein FLJ10035 (FLJ10035), mRNA
NM_030803 Homo sapiens hypothetical protein FLJ10035 (FLJ10035), mRNA NM_024062 Homo sapiens hypothetical protein MGC5338 (MGC5338), mRNA
NM 024062 Homo sapiens hypothetical protein MGC5338 (MGC5338), mRNA
NM 024059 Homo sapiens hypothetical protein MGC5356 (MGC5356), mRNA
NM_016542 Homo sapiens serine/threonine protein kinase MASK (MST4), mRNA
NM_033127 Homo sapiens regucalcin gene promotor region related protein (RGPR), mRNA
NM_033128 Homo sapiens scinderin (SCIN), mRNA
NM_033058 Homo sapiens ring finger protein 29 (RNF29), mRNA
NM_033116 Homo sapiens hypothetical protein MGC16714 (MGC16714), mRNA
NM_033123 Homo sapiens testis-development related NYD-SP27 (NYD-SP27), mRNA
NM_033126 Homo sapiens serine/threonine kinase PSKH2 (PSKH2), mRNA
NM_033124 Homo sapiens NYD-SP28 protein (NYD-SP28), mRNA
NM_033122 Homo sapiens testis development protein NYD-SP26 (NYD-SP26), mRNA
NM_033114 Homo sapiens MADP-1 protein (MADP-1), mRNA
NM_033083 Homo sapiens EAF1 protein (EAF1), mRNA
NM_033087 Homo sapiens hypothetical protein FLJ14511 (FLJ14511), mRNA
NM_024512 Homo sapiens leucine-rich repeat-containing 2 (LRRC2), mRNA
NM_006029 Homo sapiens paraneoplastic antigen MA1 (PNMA1), mRNA
NM_033025 Homo sapiens hypothetical protein FLJ13511 (7h3), mRNA
NM_015169 Homo sapiens homolog of yeast ribosome biogenesis regulatory protein RRS1
(RRS1), mRNA
NM_015129 Homo sapiens septin 6 (SEP2), mRNA
NM_032838 Homo sapiens hypothetical protein FLJ14779 (FLJ14779), mRNA
NM_032206 Homo sapiens hypothetical protein FLJ21709 (FLJ21709), mRNA
NM_032797 Homo sapiens hypothetical protein FLJ14497 (FLJ14497), mRNA
NM_032472 Homo sapiens peptidylprolyl isomerase (cyclophilin)-like 3 (PPIL3), mRNA
NM_032936 Homo sapiens DC32 (DC32), mRNA
NM_032577 Homo sapiens melanoma-associated chondroitin sulfate proteoglycan-like
(LOC84664), mRNA
NM_032933 Homo sapiens hypothetical protein MGC11386 (MGC11386), mRNA
NM_032929 Homo sapiens hypothetical protein MGC14793 (MGC14793), mRNA
NM_032928 Homo sapiens hypothetical protein MGC14141 (MGC14141), mRNA
NM 032927 Homo sapiens hypothetical protein MGC13159 (MGC13159), mRNA
NM_032926 Homo sapiens hypothetical protein MGC15737 (MGC15737), mRNA
NM_032926 Homo sapiens hypothetical protein MGC15737 (MGC15737), mRNA NM_032921 Homo sapiens hypothetical protein MGC15875 (MGC15875), mRNA
NM 032926 Homo sapiens hypothetical protein MGC15737 (MGC15737), mRNA NM 032921 Homo sapiens hypothetical protein MGC15875 (MGC15875), mRNA NM 032909 Homo sapiens hypothetical protein MGC14139 (MGC14139), mRNA
NM 032926 Homo sapiens hypothetical protein MGC15737 (MGC15737), mRNA NM 032921 Homo sapiens hypothetical protein MGC15875 (MGC15875), mRNA NM 032909 Homo sapiens hypothetical protein MGC14139 (MGC14139), mRNA NM 032908 Homo sapiens hypothetical protein MGC14407 (MGC14407), mRNA
NM 032926 Homo sapiens hypothetical protein MGC15737 (MGC15737), mRNA NM 032921 Homo sapiens hypothetical protein MGC15875 (MGC15875), mRNA NM 032909 Homo sapiens hypothetical protein MGC14139 (MGC14139), mRNA NM 032908 Homo sapiens hypothetical protein MGC14407 (MGC14407), mRNA NM 032906 Homo sapiens hypothetical protein MGC14156 (MGC14156), mRNA
NM 032926 Homo sapiens hypothetical protein MGC15737 (MGC15737), mRNA NM 032921 Homo sapiens hypothetical protein MGC15875 (MGC15875), mRNA NM 032909 Homo sapiens hypothetical protein MGC14139 (MGC14139), mRNA NM 032908 Homo sapiens hypothetical protein MGC14407 (MGC14407), mRNA

NM_032902	YT
11111_032902	Protein phosphatase 1, regulatory (inhibitor) subunit 16 A
NM 032901	(III IKIOA), IIKIVA
NM_032899	Homo sapiens hypothetical protein MGC14288 (MGC14288), mRNA
NM 032898	Homo sapiens hypothetical protein MGC14128 (MGC14128), mRNA
NM 032897	Homo sapiens hypothetical protein MGC14126 (MGC14126), mRNA
NM 032896	Homo sapiens hypothetical protein MGC14436 (MGC14436), mRNA
NM 032892	nonio sapiens hypothetical protein MGC14388 (MGC14388) mpNA
NM 032891	Tionio sapiens hypothetical protein MGC14161 (MGC14161) mDNA
NM_032890	Tiomo sapiens hypothetical protein MGC12928 (MGC12928) mDNA
NM_032887	Tionio sapiens hypothetical protein MGC13130 (MGC13130) mpN/A
NM 032885	Homo sapiens hypothetical protein MGC16037 (MGC16037), mRNA
NM 032882	Homo sapiens hypothetical protein MGC15906 (MGC15906), mRNA
NM_032881	Homo sapiens hypothetical protein MGC15827 (MGC15827), mRNA
NM_032880	Tiomo sapiens U/ shkip-specific Sm-like protein I SM10 (I SM10) DNIA
NM_032878	Tiomo sapiens hypothetical protein MGC15/30/ (MGC15/30) mDNA
NM_032873	Homo sapiens hypothetical protein MGC15677 (MGC15677), mRNA
NM 032867	Homo sapiens hypothetical protein MGC15437 (MGC15437), mRNA
NM_032865	Homo sapiens hypothetical protein FLJ14966 (FLJ14966), mRNA
NM_032861	Homo sapiens hypothetical protein FLJ14950 (FLJ14950), mRNA
NM_032859	Homo sapiens hypothetical protein FLJ14917 (FLJ14917), mRNA
NM_032856	Homo sapiens hypothetical protein FLJ14906 (FLJ14906), mRNA
NM_032855	Homo sapiens hypothetical protein FLJ14888 (FLJ14888), mRNA
NM 032854	Homo sapiens hematopoietic SH2 protein (HSH2), mRNA
NM_032850	Homo sapiens hypothetical protein FLJ14871 (FLJ14871), mRNA
NM_032849	Homo sapiens hypothetical protein FLJ14840 (FLJ14840), mRNA
NM_032847	Homo sapiens hypothetical protein FLJ14834 (FLJ14834), mRNA
NM_032846	Homo sapiens hypothetical protein FLJ14825 (FLJ14825), mRNA
NM 032844	Homo sapiens hypothetical protein FLJ14824 (FLJ14824), mRNA
NM_032843	Homo sapiens hypothetical protein FLJ14813 (FLJ14813), mRNA Homo sapiens hypothetical protein FLJ14813 (FLJ14813)
NM_032842	Homo sapiens hypothetical protein FLJ14810 (FLJ14810), mRNA Homo sapiens hypothetical protein FLJ14803 (FLJ14803), mRNA
NM_032840	Homo sapiens hypothetical protein FLJ14803 (FLJ14803), mRNA  Homo sapiens hypothetical protein FLJ14800 (FLJ14800), mRNA
NM_032839	Homo sapiens hypothetical protein FLJ14784 (FLJ14784), mRNA  Homo sapiens hypothetical protein FLJ14784 (FLJ14784), mRNA
NM_032837	Homo sapiens hypothetical protein FLJ14784 (FLJ14784), mRNA Homo sapiens hypothetical protein FLJ14775 (FLJ14775), mRNA
NM_032836	Homo sapiens hypothetical protein FLJ14768 (FLJ14768), mRNA  Homo sapiens hypothetical protein FLJ14768 (FLJ14768), mRNA
NM_032834	Homo sapiens hypothetical protein FLJ14768 (FLJ14768), mRNA  Homo sapiens hypothetical protein FLJ14751 (FLJ14751), mRNA
NM_032833	Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 15B
	(PPP1R15B), mRNA
NM_032832	Homo sapiens GARL: III protein FLJ14735 (FLJ14735), mRNA
NM_032831	Homo sapiens CAP-binding protein complex interacting protein 2 (CBCIP2),
	IMMAA
NM_032830	Homo sapiens hypothetical protein FLJ14728 (FLJ14728), mRNA
NM_032829	Homo sapiens hypothetical protein FI 114721 (FI 114721) DNIA
NM_032828	Tiolio sapiens ubiquitin UBF-fl (UBF-fl) mRNA
NM_032827	Homo Sapiens hypothetical protein FLI14708 (FLI14708) mpala
NM_032826	from sapiens hypothetical protein FLI14697 (FLI14697) mpNIA
NM_032825	Hollio sapiens hypothetical protein FI 114686 (FI 114686) mpN/A
NM_032821	Tiomo sapiens hypothetical protein FLI14665 (FLI14665) mPNIA
NM_032817	Homo sapiens hypothetical protein FI 114641 (FI 114641) mpNIA
NM_032816	fiolio sapielis hypothetical protein FLI14640 (FLI14640) mPNIA
NM_032814	fromo sapiens hypothetical protein FLI14627 (FLI14627) mRNA
NM_032811	Homo sapiens hypothetical protein FLJ14621 (FLJ14621), mRNA
	(

NM 032810	Homo sapiens hypothetical protein FLJ14600 (FLJ14600), mRNA
NM 032809	Homo sapiens hypothetical protein FLJ14596 (FLJ14596), mRNA
NM 032808	Homo sapiens hypothetical protein FLJ14594 (FLJ14594), mRNA
NM 032807	Homo sapiens hypothetical protein FLJ14590 (FLJ14590), mRNA
NM 032806	Homo sapiens hypothetical protein FLJ14566 (FLJ14566), mRNA
NM 032805	Homo sapiens hypothetical protein FLJ14549 (FLJ14549), mRNA
NM 032802	Homo sapiens hypothetical protein FLJ14540 (FLJ14540), mRNA
NM 032799	Homo sapiens hypothetical protein FLJ14524 (FLJ14524), mRNA
NM 032796	Homo sapiens reserved (SYAP1), mRNA
NM 032792	Homo sapiens hypothetical protein FLJ14486 (FLJ14486), mRNA
NM 032790	Homo sapiens hypothetical protein FLJ14466 (FLJ14466), mRNA
NM 032788	Homo sapiens hypothetical protein FLJ14457 (FLJ14457), mRNA
NM 032787	Homo sapiens hypothetical protein FLJ14454 (FLJ14454), mRNA
NM 032786	Homo sapiens hypothetical protein FLJ14451 (FLJ14451), mRNA
NM 032785	Homo sapiens hypothetical protein FLJ14442 (FLJ14442), mRNA
NM 032781	Homo sapiens hypothetical protein FLJ14427 (FLJ14427), mRNA
<del></del>	
NM_032780	Homo sapiens hypothetical protein FLJ14399 (FLJ14399), mRNA  Homo sapiens hypothetical protein FLJ14397 (FLJ14397), mRNA
NM_032779	
NM_032778	Homo sapiens hypothetical protein FLJ14393 (FLJ14393), mRNA Homo sapiens hypothetical protein FLJ14360 (FLJ14360), mRNA
NM_032775	
NM_032773	Homo sapiens hypothetical protein MGC4126 (MGC4126), mRNA
NM_032772	Homo sapiens hypothetical protein MGC2555 (MGC2555), mRNA
NM_032771	Homo sapiens hypothetical protein MGC12217 (MGC12217), mRNA
NM_032770	Homo sapiens hypothetical protein MGC16291 (MGC16291), mRNA
NM_032765	Homo sapiens hypothetical protein MGC16175 (MGC16175), mRNA
NM_032764	Homo sapiens hypothetical protein MGC16153 (MGC16153), mRNA
NM_032762	Homo sapiens hypothetical protein MGC16121 (MGC16121), mRNA
NM_032761	Homo sapiens hypothetical protein MGC16075 (MGC16075), mRNA
NM_032759	Homo sapiens hypothetical protein FLJ11328 (FLJ11328), mRNA
NM_032758	Homo sapiens hypothetical protein MGC1346 (MGC1346), mRNA
NM_032757	Homo sapiens hypothetical protein MGC15705 (MGC15705), mRNA
NM_032755	Homo sapiens hypothetical protein MGC15634 (MGC15634), mRNA
NM_032751	Homo sapiens hypothetical protein MGC15504 (MGC15504), mRNA
NM_032750	Homo sapiens hypothetical protein MGC15429 (MGC15429), mRNA
NM_032747	Homo sapiens hypothetical protein MGC14697 (MGC14697), mRNA
NM_032746	Homo sapiens hypothetical protein MGC12538 (MGC12538), mRNA
NM_032740	Homo sapiens hypothetical protein MGC5391 (MGC5391), mRNA
NM_032739	Homo sapiens hypothetical protein MGC5370 (MGC5370), mRNA
NM_032735	Homo sapiens hypothetical protein MGC13168 (MGC13168), mRNA
NM_032733	Homo sapiens hypothetical protein MGC12679 (MGC12679), mRNA
NM_032732	Homo sapiens hypothetical protein MGC10763 (MGC10763), mRNA
NM_032731	Homo sapiens hypothetical protein MGC14353 (MGC14353), mRNA
NM_032730	Homo sapiens NOGO-interacting mitochondrial protein (NIMP), mRNA
NM_032727	Homo sapiens internexin neuronal intermediate filament protein, alpha (INA),
	mRNA
NM_032726	Homo sapiens hypothetical protein MGC12837 (MGC12837), mRNA
NM_032725	Homo sapiens hypothetical protein MGC13125 (MGC13125), mRNA
NM_032724	Homo sapiens hypothetical protein MGC13269 (MGC13269), mRNA
NM_032722	Homo sapiens hypothetical protein MGC13275 (MGC13275), mRNA
NM_032721	Homo sapiens hypothetical protein MGC11314 (MGC11314), mRNA
NM_032718	Homo sapiens hypothetical protein MGC11332 (MGC11332), mRNA

ND ( 000515	
NM_032717	Homo sapiens hypothetical protein MGC11324 (MGC11324), mRNA
NM_032714	Homo sapiens hypothetical protein MGC13251 (MGC13251), mRNA
NM_032710	Homo sapiens hypothetical protein MGC13053 (MGC13053), mRNA
NM_032709	Homo sapiens hypothetical protein MGC13047 (MGC13047), mRNA
NM_032701	Homo sapiens hypothetical protein MGC2705 (MGC2705), mRNA
NM_032691	Homo sapiens hypothetical protein MGC11082 (MGC11082), mRNA
NM_032690	Homo sapiens hypothetical protein MGC13198 (MGC13198), mRNA
NM_032687	Homo sapiens hypothetical protein MGC13010 (MGC13010), mRNA
NM_032683	Homo sapiens hypothetical protein MGC12972 (MGC12972), mRNA
NM_032680	Homo sapiens hypothetical protein MGC4266 (MGC4266), mRNA
NM_032679	Homo sapiens hypothetical protein MGC4400 (MGC4400), mRNA
NM_032676	Homo sapiens hypothetical protein MGC10955 (MGC10955), mRNA
NM_032673	Homo sapiens hypothetical protein MGC10882 (MGC10882), mRNA
NM_032671	Homo sapiens hypothetical protein MGC10814 (MGC10814), mRNA
NM_032664	Homo sapiens hypothetical protein MGC11141 (MGC11141), mRNA
NM 032663	Homo sapiens hypothetical protein MGC10702 (MGC10702), mRNA
NM_032658	Homo sapiens hypothetical protein MGC10701 (MGC10701), mRNA
NM_032654	Homo sapiens hypothetical protein MGC10981 (MGC10981), mRNA
NM_032653	Homo sapiens hypothetical protein MGC10960 (MGC10960), mRNA
NM_032648	Homo sapiens hypothetical protein MGC10820 (MGC10820), mRNA
NM_032647	Homo sapiens hypothetical protein MGC10561 (MGC10561), mRNA
NM_032644	Homo sapiens hypothetical protein MGC2452 (MGC2452), mRNA
NM_032641	Homo sapiens hypothetical protein MGC2519 (MGC2519), mRNA
NM_032638	Homo sapiens hypothetical protein MGC2306 (MGC2306), mRNA
NM_032633	Homo sapiens hypothetical protein MGC5457 (MGC5457), mRNA
NM_032632	Homo sapiens hypothetical protein MGC5378 (MGC5378), mRNA
NM_032630	Homo sapiens HeLa cyclin-dependent kinase 2 interacting protein (CINP),
ND 6 022627	mRNA
NM_032627	Homo sapiens hypothetical protein MGC3181 (MGC3181), mRNA
NM_032626	Homo sapiens hypothetical brain protein my038 (MY038), mRNA
NM_032624	Homo sapiens hypothetical brain protein my050 (MY050), mRNA
NM_032623	Homo sapiens ovary-specific acidic protein (OSAP), mRNA
NM_032622	Homo sapiens multi-PDZ-domain-containing protein (LNX), mRNA
NM_032620	Homo sapiens mitochondrial GTP binding protein (GTPBG3), mRNA
NM_018622	Homo sapiens presenilins associated rhomboid-like protein (PARL), mRNA
NM_032498	Homo sapiens homeobox protein from AL590526 (LOC84528), mRNA
NM_032600	Homo sapiens testes development-related NYD-SP17 (NYD-SP17), mRNA
NM_032599	Homo sapiens testes development-related NYD-SP18 (NYD-SP18), mRNA
NM 032594	Homo sapiens insulinoma-associated protein IA-6 (INSM2), mRNA
NM_032585	Homo sapiens testis-specific transcript, Y-linked 6 (TTTY6), mRNA
NM 032575	Homo sapiens Kruppel-like zinc finger protein GLIS2 (GLIS2), mRNA
NM_032573	Homo sapiens testis-specific protein TSP-NY (TSP-NY), mRNA
NM_032572	Homo sapiens ribonuclease 7 (RNASE7), mRNA
NM_032568	Homo sapiens GABA(A) receptors associated protein like 3 (GABARAPL3),
NIM 022567	mRNA
NM_032567	Homo sapiens testis-specific protein NYD-TSP1 (NYD-TSP1), mRNA
NM_032566	Homo sapiens esophagus cancer-related gene-2 (ECG2), mRNA
NM_032562	Homo sapiens group XIII secreted phospholipase A2 (PLA2G13), mRNA
NM_032547	Homo sapiens short coiled-coil protein (HRIHFB2072), mRNA
NM_032546	Homo sapiens ring finger protein 30 (RNF30), mRNA
NM_032519 NM_032513	Homo sapiens hypothetical protein HT023 (HT023), mRNA Homo sapiens hypothetical protein MGC11303 similar to Zink transporter 2

	Larger 1999) Pari
ND ( 022400	(MGC11303), mRNA
NM_032490	Homo sapiens PNAS-127 protein (PNAS-127), mRNA
NM_032488	Homo sapiens protein related with psoriasis (LOC84518), mRNA
NM_032471	Homo sapiens protein kinase (cAMP-dependent, catalytic) inhibitor beta (PKIB), mRNA
NM_032292	Homo sapiens hypothetical protein FLJ20203 (FLJ20203), mRNA
NM_032263	Homo sapiens hypothetical protein DKFZp434B227 (DKFZp434B227), mRNA
NM 015178	Homo sapiens KIAA0717 protein (KIAA0717), mRNA
NM_032410	Homo sapiens hook3 protein (HOOK3), mRNA
NM_032108	Homo sapiens sema domain, transmembrane domain (TM), and cytoplasmic
	domain, (semaphorin) 6B (SEMA6B), mRNA
NM_015636	Homo sapiens DKFZP586J0119 protein (DKFZP586J0119), mRNA
NM_015701	Homo sapiens hypothetical protein (CL25084), mRNA
NM_015224	Homo sapiens KIAA1105 protein (RAP140), mRNA
NM_032390	Homo sapiens nucleolar protein interacting with the FHA domain of pKi-67 (NIFK), mRNA
NM_032388	Homo sapiens nasopharyngeal carcinoma-related protein (NPCR), mRNA
NM_032383	Homo sapiens Hermansky-Pudlak syndrome 3 (HPS3), mRNA
NM_032378	Homo sapiens hypothetical protein FLJ20897 (FLJ20897), mRNA
NM_032376	Homo sapiens hypothetical protein MGC4251 (MGC4251), mRNA
NM_032375	Homo sapiens hypothetical protein MGC2865 (MGC2865), mRNA
NM_032373	Homo sapiens hypothetical protein MGC16202 (MGC16202), mRNA
NM_032370	Homo sapiens hypothetical protein MGC15716 (MGC15716), mRNA
NM_032369	Homo sapiens hypothetical protein MGC15619 (MGC15619), mRNA
NM_032368	Homo sapiens hypothetical protein MGC15436 (MGC15436), mRNA
NM_032374	Homo sapiens hypothetical protein MGC2562 (MGC2562), mRNA
NM_032364	Homo sapiens hypothetical protein MGC14726 (MGC14726), mRNA
NM_032362	Homo sapiens HEIL1 protein (HEIL1), mRNA
NM_032361	Homo sapiens hypothetical protein MGC5469 (MGC5469), mRNA
NM_032360	Homo sapiens hypothetical protein MGC2404 (MGC2404), mRNA
NM_032359	Homo sapiens hypothetical protein MGC4308 (MGC4308), mRNA
NM_032358	Homo sapiens hypothetical protein MGC13183 (MGC13183), mRNA
NM_032357	Homo sapiens hypothetical protein MGC12981 (MGC12981), mRNA
NM_032356	Homo sapiens hypothetical protein MGC14151 (MGC14151), mRNA
NM_032355	Homo sapiens hypothetical protein MGC13272 (MGC13272), mRNA
NM_032352	Homo sapiens hypothetical protein MGC11296 (MGC11296), mRNA
NM_032350	Homo sapiens hypothetical protein MGC11257 (MGC11257), mRNA
NM_032349	Homo sapiens hypothetical protein MGC11275 (MGC11275), mRNA
NM_032348	Homo sapiens hypothetical protein MGC3047 (MGC3047), mRNA
NM_032346	Homo sapiens hypothetical protein MGC13096 (MGC13096), mRNA
NM_032345	Homo sapiens hypothetical protein MGC13064 (MGC13064), mRNA
NM_032343	Homo sapiens hypothetical protein MGC13016 (MGC13016), mRNA
NM_032341	Homo sapiens hypothetical protein MGC14844 (MGC14844), mRNA
NM_032339	Homo sapiens hypothetical protein MGC14832 (MGC14832), mRNA
NM_032336	Homo sapiens hypothetical protein MGC14799 (MGC14799), mRNA
NM_032334	Homo sapiens hypothetical protein MGC14595 (MGC14595), mRNA
NM_032332	Homo sapiens hypothetical protein MGC4238 (MGC4238), mRNA
NM_032331	Homo sapiens hypothetical protein MGC2408 (MGC2408), mRNA
NM_032328	Homo sapiens hypothetical protein MGC12458 (MGC12458), mRNA
NM_032322	Homo sapiens hypothetical protein MGC13061 (MGC13061), mRNA
NM_032321	Homo sapiens hypothetical protein MGC13057 (MGC13057), mRNA
NM_032319	Homo sapiens chromosome 2 open reading frame 7 (C2orf7), mRNA
	1 77 411

	T
NM_032315	Homo sapiens hypothetical protein MGC4399 (MGC4399), mRNA
NM_032314	Homo sapiens hypothetical protein MGC4767 (MGC4767), mRNA
NM_032313	Homo sapiens hypothetical protein MGC3232 (MGC3232), mRNA
NM_032312	Homo sapiens hypothetical protein MGC11061 (MGC11061), mRNA
NM_032310	Homo sapiens hypothetical protein MGC11115 (MGC11115), mRNA
NM_032307	Homo sapiens hypothetical protein MGC10999 (MGC10999), mRNA
NM_032303	Homo sapiens hypothetical protein MGC10940 (MGC10940), mRNA
NM_032302	Homo sapiens hypothetical protein MGC10911 (MGC10911), mRNA
NM_032301	Homo sapiens hypothetical protein MGC10870 (MGC10870), mRNA
NM_032300	Homo sapiens hypothetical protein MGC10854 (MGC10854), mRNA
NM_032298	Homo sapiens hypothetical protein DKFZp761O132 (DKFZp761O132), mRNA
NM_032297	Homo sapiens hypothetical protein DKFZp761D112 (DKFZp761D112), mRNA
NM_032296	Homo sapiens hypothetical protein DKFZp761A132 (DKFZp761A132), mRNA
NM_032295	Homo sapiens hypothetical protein DKFZp761N0624 (DKFZp761N0624), mRNA
NM_032294	Homo sapiens hypothetical protein DKFZp761M0423 (DKFZp761M0423), mRNA
NM_032289	Homo sapiens hypothetical protein DKFZp761B0514 (DKFZp761B0514), mRNA
NM_032287	Homo sapiens hypothetical protein DKFZp761O17121 (DKFZp761O17121), mRNA
NM_032280	Homo sapiens hypothetical protein DKFZp761J139 (DKFZp761J139), mRNA
NM_032278	Homo sapiens hypothetical protein DKFZp547P082 (DKFZp547P082), mRNA
NM_032274	Homo sapiens hypothetical protein DKFZp547F072 (DKFZp547F072), mRNA
NM_032271	Homo sapiens hypothetical protein DKFZp586I021 (DKFZp586I021), mRNA
NM_032270	Homo sapiens hypothetical protein DKFZp586J1119 (DKFZp586J1119), mRNA
NM_032269	Homo sapiens hypothetical protein DKFZp434I099 (DKFZp434I099), mRNA
NM_032266	Homo sapiens hypothetical protein DKFZp434G118 (DKFZp434G118), mRNA
NM_032265	Homo sapiens hypothetical protein DKFZp434N127 (DKFZp434N127), mRNA
NM_032262	Homo sapiens hypothetical protein DKFZp434N035 (DKFZp434N035), mRNA
NM_032257	Homo sapiens hypothetical protein DKFZp434N2435 (DKFZp434N2435), mRNA
NM_032256	Homo sapiens hypothetical protein DKFZp434K2435 (DKFZp434K2435), mRNA
NM_032255	Homo sapiens hypothetical protein DKFZp434I1930 (DKFZp434I1930), mRNA
NM_032254	Homo sapiens hypothetical protein DKFZp434F142 (DKFZp434F142), mRNA
NM_032247	Homo sapiens hypothetical protein DKFZp434E0519 (DKFZp434E0519), mRNA
NM_032242	Homo sapiens hypothetical protein DKFZp564A176 (DKFZp564A176), mRNA
NM_032238	Homo sapiens hypothetical protein FLJ23416 (FLJ23416), mRNA
NM_032235	Homo sapiens hypothetical protein FLJ23138 (FLJ23138), mRNA
NM_032234	Homo sapiens hypothetical protein FLJ23059 (FLJ23059), mRNA
NM_032233	Homo sapiens hypothetical protein FLJ23027 (FLJ23027), mRNA
NM_032229	Homo sapiens hypothetical protein FLJ22774 (FLJ22774), mRNA
NM_032221	Homo sapiens hypothetical protein FLJ22369 (FLJ22369), mRNA
NM_032213	Homo sapiens hypothetical protein FLJ21977 (FLJ21977), mRNA
NM_032212	Homo sapiens similar to DNA-directed RNA polymerase I (135 kDa) (Rpo1-2), mRNA
NM_032207	Homo sapiens hypothetical protein FLJ21742 (FLJ21742), mRNA
NM 032205	Homo sapiens hypothetical protein FLJ21615 (FLJ21615), mRNA
NM 032196	Homo sapiens hypothetical protein KIAA1259 (KIAA1259), mRNA
NM 032192	Homo sapiens hypothetical protein FLJ20940 (FLJ20940), mRNA
	captons hypothetical protein i 1320740 (PL320740), HIRINA

NM_032191	Homo sapiens hypothetical protein FLJ14326 (FLJ14326), mRNA
NM_032187	Homo sapiens hypothetical protein FLJ14026 (FLJ14026), mRNA
NM_032186	Homo sapiens hypothetical protein FLJ13964 (FLJ13964), mRNA
NM_032181	Homo sapiens hypothetical protein FLJ13391 (FLJ13391), mRNA
NM_032179	Homo sapiens hypothetical protein FLJ20542 (FLJ20542), mRNA
NM_032178	Homo sapiens hypothetical protein FLJ13291 (FLJ13291), mRNA
NM 032175	Homo sapiens hypothetical protein FLJ12787 (FLJ12787), mRNA
NM 032174	Homo sapiens hypothetical protein FLJ12770 (FLJ12770), mRNA
NM 032169	Homo sapiens hypothetical protein FLJ12592 (FLJ12592), mRNA
NM_032164	Homo sapiens hypothetical protein FLJ12298 (FLJ12298), mRNA
NM_032162	Homo sapiens hypothetical protein FLJ11952 (FLJ11952), mRNA
NM_032155	Homo sapiens hypothetical protein DKFZp547I094 (DKFZp547I094), mRNA
NM 032152	Homo sapiens PRAM-1 protein (PRAM-1), mRNA
NM_032149	Homo sapiens hypothetical protein DKFZp434G072 (DKFZP434G072), mRNA
NM_032147	Homo sapiens hypothetical protein DKFZp434D0127 (DKFZP434D0127), mRNA
NM_032146	Homo sapiens hypothetical protein DKFZp434L1123 similar to mouse Arl6
	(DKFZP434L1123), mRNA
NM_032143	Homo sapiens hypothetical protein DKFZp434B1727 (DKFZP434B1727), mRNA
NM_032142	Homo sapiens hypothetical protein FLJ10352 (FLJ10352), mRNA
NM_032141	Homo sapiens hypothetical protein DKFZp434K1421 (DKFZP434K1421), mRNA
NM_032140	Homo sapiens hypothetical protein DKFZp434A1319 (DKFZP434A1319), mRNA
NM_032135	Homo sapiens hypothetical protein DKFZp434F1017 (DKFZP434F1017), mRNA
NM_032134	Homo sapiens hypothetical protein DKFZp434P0316 (DKFZP434P0316), mRNA
NM_032131	Homo sapiens hypothetical protein DKFZp434P0714 (DKFZP434P0714), mRNA
NM 032130	Homo sapiens hypothetical protein DKFZp434J0113 (DKFZP434J0113), mRNA
NM_032129	Homo sapiens hypothetical protein DKFZp434H2010 (DKFZP434H2010), mRNA
NM 032128	Homo sapiens hypothetical protein DKFZp566M114 (DKFZP566M114), mRNA
NM_032127	Homo sapiens hypothetical protein DKFZp566M1046 (DKFZP566M1046), mRNA
NM 032126	Homo sapiens hypothetical protein DKFZp564J047 (DKFZP564J047), mRNA
NM 032124	Homo sapiens hypothetical protein DKFZp564D1378 (DKFZP564D1378),
	mRNA
NM 032121	Homo sapiens hypothetical protein DKFZp564K142 similar to implantation-
	associated protein (DKFZp564K142), mRNA
NM 032118	Homo sapiens hypothetical protein FLJ12953 similar to Mus musculus D3Mm3e
	(FLJ12953), mRNA
NM 032117	Homo sapiens GAJ protein (GAJ), mRNA
NM_032116	Homo sapiens hypothetical protein MGC2599 similar to katanin p60 subunit A 1 2599 (MGC2599), mRNA
NM 032112	Homo sapiens mitochondrial ribosomal protein L43 (MRPL43), mRNA
NM 020898	Homo sapiens KIAA1536 protein (KIAA1536), mRNA
NM_020726	Homo sapiens neurolysin (metallopeptidase M3 family) (NLN), mRNA
NM 020707	Homo sapiens KIAA1173 protein (KIAA1173), mRNA
NM 018670	Homo sapiens hypothetical protein (IR1899308), mRNA
	1 Special Control of the Control of

NM_018385 Homo sapiens hypothetical protein FLJ11301 (FLJ11301), mRNA NM_018064 Homo sapiens hypothetical protein FLJ10342 (FLJ10342), mRNA	
<del></del>	
NM_017607 Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 12C (PPP1R12C), mRNA	
NM_015645 Homo sapiens DKFZP586B0621 protein (CTRP5), mRNA	
NM_015528 Homo sapiens DKFZP566H073 protein (DKFZP566H073), mRNA	
NM_015512 Homo sapiens DKFZP434A236 protein (DKFZP434A236), mRNA	
NM_015426 Homo sapiens DKFZP434C245 protein (DKFZP434C245), mRNA	
NM_015292 Homo sapiens KIAA0747 protein (KIAA0747), mRNA	
NM_015236 Homo sapiens KIAA0768 protein (LEC3), mRNA	
NM_015196 Homo sapiens KIAA0922 protein (KIAA0922), mRNA	
NM_015112 Homo sapiens KIAA0807 protein (MAST205), mRNA	
NM_015070 Homo sapiens KIAA0853 protein (KIAA0853), mRNA	
NM_032308 Homo sapiens hypothetical protein MGC4189 (MGC4189), mRNA	
NM_004801 Homo sapiens neurexin 1 (NRXN1), mRNA	
NM_001221 Homo sapiens calcium/calmodulin-dependent protein kinase (CaM kinase) II	
delta (CAMK2D), mRNA	
NM_015208 Homo sapiens KIAA0874 protein (KIAA0874), mRNA	
NM_032043 Homo sapiens BRCA1-interacting protein 1 (BRIP1), mRNA	
NM 032040 Homo sapiens hypothetical protein DKFZp564K0322 (DKFZP564K0322),	
mRNA	
NM_032037 Homo sapiens serine/threonine protein kinase SSTK (SSTK), mRNA	
NM_032033 Homo sapiens FKSG43 (FKSG43), mRNA	
NM_032032 Homo sapiens FKSG42 (FKSG42), mRNA	
NM_032031 Homo sapiens FKSG17 (FKSG17), mRNA	
NM_032029 Homo sapiens FKSG87 protein (FKSG87), mRNA	
NM_032026 Homo sapiens CDA11 protein (CDA11), mRNA	
NM_032024 Homo sapiens CDA017 protein (CDA017), mRNA	
NM_032023 Homo sapiens AD037 protein (AD037), mRNA	
NM_032022 Homo sapiens AD036 protein (AD036), mRNA	
NM_031956 Homo sapiens NYD-SP14 protein (NYD-SP14), mRNA	
NM_031954 Homo sapiens MSTP028 protein (MSTP028), mRNA	
NM_031953 Homo sapiens MSTP043 protein (MSTP043), mRNA	
NM_031936 Homo sapiens G protein-coupled receptor 61 (GPR61), mRNA	
NM_031934 Homo sapiens RAB34, member RAS oncogene family (RAB34), mRNA	
NM_031933 Homo sapiens wingless-type MMTV integration site family, member 8A	
(WNT8A), transcript variant 1, mRNA	
NM_031932 Homo sapiens testis transcript Y 14 (TTY14), mRNA	-
NM_031931 Homo sapiens testis transcript Y 13 (TTY13), mRNA	
NM_031930 Homo sapiens testis transcript Y 12 (TTY12), mRNA	
NM_031929 Homo sapiens testis transcript Y 11 (TTY11), mRNA	_
NM_031927 Homo sapiens testis transcript Y 9 (TTY9), mRNA	
NM_031926 Homo sapiens testis transcript Y 7 (TTY7), mRNA	
NM_031925 Homo sapiens transmembrane protein induced by tumor necrosis factor alpha	
(TMPIT), mRNA	
NM_031924 Homo sapiens radial spoke protein 3 (RSP3), mRNA	
NM_031917 Homo sapiens angiopoietin-related protein 5 (ARP5), mRNA	
NM_031948 Homo sapiens marapsin (MPN), mRNA	
NM_031908 Homo sapiens complement-c1q tumor necrosis factor-related protein 2 (CTRP2	2),
mRNA	
NM_031905 Homo sapiens hypothetical protein MGC3195 (MGC3195), mRNA	
NM_031889 Homo sapiens enamelin (ENAM), mRNA	

NM_022447	Homo sapiens topoisomerase-related function protein 4-2 (TRF4-2), mRNA
NM_031485	Homo sapiens glutamate rich WD repeat protein GRWD (GRWD), mRNA
NM_031484	Homo sapiens hypothetical protein MGC4415 (MGC4415), mRNA
NM_031479	Homo sapiens hypothetical protein MGC4638 (MGC4638), mRNA
NM_031474	Homo sapiens hypothetical protein DKFZp761G1913 (DKFZP761G1913), mRNA
NM_031466	Homo sapiens KIAA1882 protein (MGC4737), mRNA
NM_031465	Homo sapiens hypothetical protein MGC13204 (MGC13204), mRNA
NM_031464	Homo sapiens hypothetical protein MGC11287 similar to ribosomal protein S6 kinase, (MGC11287), mRNA
NM 031459	Homo sapiens sestrin 2 (SES2), mRNA
NM 031455	Homo sapiens hypothetical protein DKFZp761F241 (DKFZP761F241), mRNA
NM 031453	Homo sapiens hypothetical protein MGC11034 (MGC11034), mRNA
NM 031452	Homo sapiens hypothetical protein MGC2560 (MGC2560), mRNA
NM 031449	Homo sapiens KIAA1886 protein (DKFZP761I2123), mRNA
NM 031447	Homo sapiens hypothetical protein MGC13033 (MGC13033), mRNA
NM 031446	Homo sapiens hypothetical protein PNAS-131 (PNAS-131), mRNA
NM 031437	Homo sapiens hypothetical protein MGC10823 (MGC10823), mRNA
NM 031436	Homo sapiens hypothetical protein MGC10612 (MGC10612), mRNA
NM 031435	Homo sapiens hypothetical protein DKFZp564I0422 (DKFZP564I0422), mRNA
NM 031430	Homo sapiens rab interacting lysosomal protein (RILP), mRNA
NM 031425	Homo sapiens hypothetical protein MGC10812 (MGC10812), mRNA
NM 031423	Homo sapiens hypothetical protein NUF2R (NUF2R), mRNA
NM_031421	Homo sapiens hypothetical protein DKFZp434H0115 (DKFZP434H0115), mRNA
NM_031412	Homo sapiens GABA(A) receptor-associated protein like 1 (GABARAPL1), mRNA
NM 004637	Homo sapiens RAB7, member RAS oncogene family (RAB7), mRNA
NM 031283	Homo sapiens HMG-box transcription factor TCF-3 (TCF-3), mRNA
NM 031307	Homo sapiens hypothetical protein FKSG32 (FKSG32), mRNA
NM_031305	Homo sapiens hypothetical protein DKFZp564B1162 (DKFZP564B1162), mRNA
NM_031301	Homo sapiens hypothetical protein DKFZp564D0372 (DKFZP564D0372), mRNA
NM_031298	Homo sapiens hypothetical protein MGC2963 (MGC2963), mRNA
NM_031293	Homo sapiens hypothetical protein DKFZp434G131 (DKFZP434G131), mRNA
NM_031292	Homo sapiens hypothetical protein DKFZp434G1415 (DKFZP434G1415), mRNA
NM_031288	Homo sapiens PAP-1 binding protein (PAPA-1), mRNA
NM_031284	Homo sapiens hypothetical protein DKFZp434B195 (DKFZP434B195), mRNA
NM 030972	Homo sapiens hypothetical protein MGC5384 (MGC5384), mRNA
NM_030901	Homo sapiens olfactory receptor, family 7, subfamily A, member 17 (OR7A17), mRNA
NM_017990	Homo sapiens hypothetical protein FLJ10079 (FLJ10079), mRNA
NM_031219	Homo sapiens hypothetical protein MGC12904 (MGC12904), mRNA
NM 031218	Homo sapiens hypothetical protein FLJ12488 (FLJ12488), mRNA
NM 031214	Homo sapiens hypothetical protein AF311304 (AF311304), mRNA
NM 031210	Homo sapiens hypothetical protein DC50 (DC50), mRNA
NM 031207	Homo sapiens hypothetical protein HT036 (HT036), mRNA
NM 007013	Homo sapiens WW domain-containing protein 1 (WWP1), mRNA
NM 030897	Homo sapiens hypothetical protein FLJ21617 (FLJ21617), mRNA
NM 030978	Homo sapiens hypothetical protein similar to actin related protein 2/3 complex,

Г <del></del>	
ND 4 000071	subunit 5 (MGC3038), mRNA
NM_030971	Homo sapiens similar to rat tricarboxylate carrier-like protein (BA108L7.2),
77.5	IIICIVA
NM_030965	Homo sapiens similar to sialyltransferase 7 ((alpha-N-acetylneuraminyl 2,3-
	Total galactosyl-1,3/-in-acetyl galactosaminide alpha-2 6-sialyltransforms T
	(MGC3164), IRNA
NM_030960	Homo sapiens sperm acrosome associated 1 (SPACA1), mRNA
NM_030958	nomo sapiens organic anion transporter polypentide-related protein 4
	(OATERF4), mrnA
NM_030952	Homo sapiens hypothetical protein DKF7p4341037 (DKF7p4247037)
NM_030940	Homo sapiens hypothetical protein MGC4276 similar to CG8198 (MGC4276),
NM_030937	Homo sapiens hypothetical protein hCLA-iso (HCLA-ISO), mRNA
NM_030929	Homo sapiens hypothetical protein FKSG28 (FKSG28), mRNA
NM_030921	Homo sapiens hypothetical protein DC42 (DC42), mRNA
NM_030917	Homo sapiens hypothetical protein DKFZp586K0717 (DKFZP586K0717),
_	mRNA
NM_030915	
NM_030914	Homo sapiens hypothetical protein DKFZp566J091 (DKFZP566J091), mRNA
NM 030907	Homo sapiens hypothetical protein MGC2668 (MGC2668), mRNA
NM_030895	Homo sapiens hypothetical protein MGC10731 (MGC10731), mRNA
NM_030891	Homo sapiens hypothetical protein FLJ14129 (FLJ14129), mRNA
NM_030755	Homo sapiens leucine-rich repeat-containing 3 (LRRC3), mRNA
NM_030819	Homo sapiens thioredoxin domain-containing (TXNDC), mRNA
NM 030814	110110 sapiens hypothetical protein MGC11335 (MGC11335) DATA
NM 030810	Homo sapiens hypothetical protein GL012 (GL012), mRNA
NM_030804	Hollo sapiens hypothetical protein MGC3178 (MGC3179) DNA
14141_030804	Homo sapiens hypothetical protein DKFZp434F2135 (DKFZp434F2135)
NM_030794	IIIIIII
NM_030759	Homo sapiens hypothetical protein FLJ21007 (FLJ21007), mRNA
NM_030795	Homo sapiens nuclear receptor binding factor-2 (NRRE-2), mPNA
	Tiomo sapiens statifinin-like 4 (STMNA) mRNA
NM_020909	Homo sapiens KIAA1548 protein (KIAA1548) mPNIA
NM_018023	nomo sapiens hypothetical protein FI 110201 (FI 110201) DALA
NM_023009	Tiolio sapielis macrophage myristovlated alanine rich Claimannia
T. 6. 0.0	(WITCHFACES), IIRNA
VM_025230	Homo sapiens hypthetical protein PRO2389 (PRO2389), mRNA
VM_025222	Homo sapiens hypothetical protein PRO2730 (PRO2730) mPNA
VM_025170	nomo sapiens hypothetical protein FI I12987 (FI I12987) mDNIA
111 024001	Homo sapiens hypothetical protein FL I12242 (FL I12242) mpala
1212 02 7720	Tionio sapiens hypothetical protein FI 122550 (FI 122550) DATA
M_017578	Homo sapiens AKAP-binding sperm protein ropporin (DKFZp434B1222),
	muui
IM_030642	Homo sapiens apolipoprotein L, 5 (APOL5), mRNA
1111_024313	
M 4 020 CO1	Homo sapiens FYVE and coiled-coil domain containing 1 (FVCO1)
IM_030621	Homo sapiens FYVE and coiled-coil domain containing 1 (FVCO1) PNIA
050021	Homo sapiens FYVE and coiled-coil domain containing 1 (FYCO1), mRNA Homo sapiens helicase-moi (KIAA0928), mRNA
M_030641	Homo sapiens FYVE and coiled-coil domain containing 1 (FYCO1), mRNA Homo sapiens helicase-moi (KIAA0928), mRNA Homo sapiens apolipoprotein L. 6 (APOL 6), mRNA
M_030641 M_025190	Homo sapiens FYVE and coiled-coil domain containing 1 (FYCO1), mRNA Homo sapiens helicase-moi (KIAA0928), mRNA Homo sapiens apolipoprotein L, 6 (APOL6), mRNA Homo sapiens KIAA1641 protein (KIAA1641), mRNA
M 025190 M 025040	Homo sapiens FYVE and coiled-coil domain containing 1 (FYCO1), mRNA Homo sapiens helicase-moi (KIAA0928), mRNA Homo sapiens apolipoprotein L, 6 (APOL6), mRNA Homo sapiens KIAA1641 protein (KIAA1641), mRNA Homo sapiens hypothetical protein FLI21941 (FLI21941), mRNA
M 030641 M 025190 M 025040 M 030613	Homo sapiens FYVE and coiled-coil domain containing 1 (FYCO1), mRNA Homo sapiens helicase-moi (KIAA0928), mRNA Homo sapiens apolipoprotein L, 6 (APOL6), mRNA Homo sapiens KIAA1641 protein (KIAA1641), mRNA Homo sapiens hypothetical protein FLJ21941 (FLJ21941), mRNA Homo sapiens hypothetical protein FLJ21628 (FLJ21628) mRNA
M 030641 M 025190 M 025040 M 030613 M 024820	Homo sapiens FYVE and coiled-coil domain containing 1 (FYCO1), mRNA Homo sapiens helicase-moi (KIAA0928), mRNA Homo sapiens apolipoprotein L, 6 (APOL6), mRNA Homo sapiens KIAA1641 protein (KIAA1641), mRNA Homo sapiens hypothetical protein FLJ21941 (FLJ21941), mRNA Homo sapiens hypothetical protein FLJ21628 (FLJ21628), mRNA Homo sapiens KIAA1608 protein (KIAA1608), mRNA
M 030641 M 025190 M 025040 M 030613 M 024820 M 018015 I	Homo sapiens FYVE and coiled-coil domain containing 1 (FYCO1), mRNA Homo sapiens helicase-moi (KIAA0928), mRNA Homo sapiens apolipoprotein L, 6 (APOL6), mRNA Homo sapiens KIAA1641 protein (KIAA1641), mRNA Homo sapiens hypothetical protein FLJ21941 (FLJ21941), mRNA Homo sapiens hypothetical protein FLJ21628 (FLJ21628) mRNA

NT 6 00 1000	TI MCCA2A2 (MCCA2A2) - DNA
NM_024329	Homo sapiens hypothetical protein MGC4342 (MGC4342), mRNA
NM_024087	Homo sapiens DKFZP564L0862 protein (DKFZP564L0862), mRNA
NM_030594	Homo sapiens cytoplasmic polyadenylation element binding protein (CPEB1),
NB 4 025004	mRNA
NM_025084	Homo sapiens hypothetical protein FLJ22795 (FLJ22795), mRNA
NM_025090	Homo sapiens KIAA1453 protein (KIAA1453), mRNA
NM_024939	Homo sapiens hypothetical protein FLJ21918 (FLJ21918), mRNA
NM_024903	Homo sapiens hypothetical protein FLJ14297 (FLJ14297), mRNA
NM_024793	Homo sapiens KIAA0643 protein (KIAA0643), mRNA
NM_024718	Homo sapiens hypothetical protein FLJ10101 (FLJ10101), mRNA  Homo sapiens DKFZP564P1916 protein (DKFZP564P1916), mRNA
NM_015652	Homo sapiens DRFZF364F1916 protein (DRFZF364F1916), filktva  Homo sapiens hypothetical protein FLJ13659 (FLJ13659), mRNA
NM_025189	
NM_025021	Homo sapiens KIAA0616 protein (KIAA0616), mRNA
NM_025010	Homo sapiens KIAA0795 protein (KIAA0795), mRNA
NM_024894	Homo sapiens hypothetical protein FLJ14075 (FLJ14075), mRNA
NM_024840	Homo sapiens hypothetical protein FLJ13590 (FLJ13590), mRNA
NM_022782	Homo sapiens M-phase phosphoprotein 9 (MPHOSPH9), mRNA
NM_017558	Homo sapiens hypothetical protein DKFZp434L0850 (DKFZp434L0850),
NTM 020590	mRNA  Home conjugate hypothetical protein MGC10520 (MGC10520) mRNA
NM 030580	Homo sapiens hypothetical protein MGC10520 (MGC10520), mRNA Homo sapiens phosphoprotein regulated by mitogenic pathways (C8FW), mRNA
NM_025195	
NM_030581	Homo sapiens hypothetical protein FLJ12270 (FLJ12270), mRNA
NM_030577	Homo sapiens hypothetical protein MGC10993 (MGC10993), mRNA
NM_030576	Homo sapiens hypothetical protein MGC10986 (MGC10986), mRNA
NM_030575	Homo sapiens hypothetical protein MGC10334 (MGC10334), mRNA
NM_030572	Homo sapiens hypothetical protein MGC10946 (MGC10946), mRNA
NM_030571	Homo sapiens hypothetical protein MGC10924 similar to Nedd4 WW-binding protein 5 (MGC10924), mRNA
NM 030569	Homo sapiens hypothetical protein MGC10848 (MGC10848), mRNA
NM 030568	Homo sapiens hypothetical protein MGC10848 (MGC10848), mRNA
NM 030567	Homo sapiens hypothetical protein MGC10818 (MGC10818), iniciva
NM 025164	Homo sapiens KIAA0999 protein (KIAA0999), mRNA
NM 025132	Homo sapiens KIAA1638 protein (KIAA1638), mRNA
NM 024668	Homo sapiens kiaarosa protein (kiaarosa), inkina Homo sapiens hypothetical protein FLJ20288 (FLJ20288), mRNA
NM 024547	Homo sapiens KIAA0467 protein (KIAA0467), mRNA
NM 018418	Homo sapiens kypothetical protein (HSD-3.1), mRNA
	Homo sapiens hypothetical protein (H3D-3.1), filking Homo sapiens hypothetical protein FLJ11560 (FLJ11560), mRNA
NM 025182	Homo sapiens LAP (leucine-rich repeats and PDZ) and no PDZ protein (LANO),
NM_025168	mRNA
NM 025081	Homo sapiens KIAA1305 protein (KIAA1305), mRNA
NM 024750	Homo sapiens RIAA1303 protein (RIAA1303), inkiva  Homo sapiens leucine-rich repeat-containing 2 (LRRC2), mRNA
NM 025266	Homo sapiens hypothetical protein MGC2780 (MGC2780), mRNA
NM 025265	Homo sapiens hypothetical protein MGC2780 (MGC2780), mRNA  Homo sapiens hypothetical protein MGC2776 (MGC2776), mRNA
NM 025264	Homo sapiens hypothetical protein MGC2476 (MGC2476), mRNA  Homo sapiens hypothetical protein MGC2454 (MGC2454), mRNA
NM 025247	Homo sapiens hypothetical protein MGC2434 (MGC2434), mRNA  Homo sapiens hypothetical protein MGC5601 (MGC5601), mRNA
NM_025247 NM_025246	Homo sapiens hypothetical protein MGC3001 (MGC3001), mRNA  Homo sapiens hypothetical protein MGC3295 (MGC3295), mRNA
NM 025234	Homo sapiens recombination protein REC14 (REC14), mRNA
NM_025221	Homo sapiens calsenilin-like protein (CALP), mRNA
NM_025207	Homo sapiens hypothetical protein PP591 (PP591), mRNA
NM 025204	Homo sapiens hypothetical protein PP2447 (PP2447), mRNA
NM_025203	Homo sapiens hypothetical protein FLJ21945 (FLJ21945), mRNA
NM_025199	Homo sapiens hypothetical protein FLJ20886 (FLJ20886), mRNA

NM_025197 NM_025187 NM_025184 NM_025181	Homo sapiens hypothetical protein FLJ13660 similar to CDK5 activator-binding protein C53 (FLJ13660), mRNA
NM_025184	Diotem C55 (FLJ13000), mRNA
NM_025184	Homo contone le de la
	Homo sapiens hypothetical protein FLJ12076 (FLJ12076), mRNA
INIM   025   K	Homo sapiens hypothetical protein FLJ22843 (FLJ22843) mRNA
	Homo sapiens hypothetical protein FLJ22004 (FLJ22004) mPNA
NM_025163	Homo sapiens hypothetical protein FLJ12768 (FLJ12768) mPNA
NM_025159	Homo sapiens hypothetical protein FLJ11577 (FLJ11577) mPNIA
NM_025157	Homo sapiens hypothetical protein FLI23042 (FLI23042) mDNA
NM_025155	Homo sapiens hypothetical protein FLJ11848 (FLJ11848) mDNA
NM_025152	FIGURE Sapiens hypothetical protein FLI12660 (FLI12660) mPNIA
NM_025150	Homo sapiens hypothetical protein FLI12528 (FLI12528) mDNIA
NM_025147	Fiorno sapiens hypothetical protein FLI13448 (FI 113448) mDNIA
NM_025146	110110 sapiens hypothetical protein FL 113194 (FL 113194) mDNA
NM_025145	Homo sapiens hypothetical protein FLJ22944 (FLJ22944), mRNA
NM_025143	Homo sapiens hypothetical protein FLJ20856 (FLJ20856), mRNA
NM_025140	Homo sapiens hypothetical protein FLJ22471 (FLJ22471), mRNA
NM_025139	Homo sapiens hypothetical protein FLJ12584 (FLJ12584), mRNA
NM_025134	Homo sapiens hypothetical protein FLJ12178 (FLJ12178), mRNA
NM_025133	Homo sapiens hypothetical protein FLJ12673 (FLJ12673), mRNA
NM_025130	Homo sapiens hypothetical protein FLJ22761 (FLJ22761), mRNA
NM_025129	Homo sapiens hypothetical protein FLJ22688 (FLJ22688), mRNA
NM_025118	Homo sapiens hypothetical protein FLJ13310 (FLJ13310), mRNA
NM_025115	Homo sapiens hypothetical protein FLJ23263 (FLJ23263), mRNA
NM 025113	Homo saniens hypothetical protein FL 1215 (2) (FLJ23263), mRNA
NM_025112	Homo sapiens hypothetical protein FLJ21562 (FLJ21562), mRNA Homo sapiens hypothetical protein MCC1164 (FLJ21562), mRNA
NM 025108	Homo sapiens hypothetical protein MGC11349 (MGC11349), mRNA Homo sapiens hypothetical protein FI H2000 (MGC11349), mRNA
NM_025107	Homo sapiens hypothetical protein FLJ13909 (FLJ13909), mRNA
NM 025105	Homo sapiens hypothetical protein FLJ21269 (FLJ21269), mRNA Homo sapiens hypothetical protein FLJ12409 (FLJ12409), mRNA
NM_025104	Homo sapiens hypothetical protein FLJ13087 (FLJ13087), mRNA
NM_025103	Homo sapiens capillary morphogenesis protein 1 (CMG1), mRNA
NM_025100	Homo sapiens hypothetical protein FLJ12294 (FLJ12294), mRNA
NM_025093	Homo saniens hypothetical protein FL 11227 (FL 112294), mRNA
NM_025092	Homo sapiens hypothetical protein FLJ11827 (FLJ11827), mRNA  Homo sapiens hypothetical protein FLJ12237 (FLJ11827), mRNA
NM_025088	Homo sapiens hypothetical protein FLJ22635 (FLJ22635), mRNA
NM 025087	Homo sapiens hypothetical protein FLJ13241 (FLJ13241), mRNA
NM 025082	Homo sapiens hypothetical protein FLJ21511 (FLJ21511), mRNA
NM_025075	Homo sapiens hypothetical protein FLJ13111 (FLJ13111), mRNA
NM 025074	Homo sapiens hypothetical protein FLJ23445 (FLJ23445), mRNA
NM_025073	Homo sapiens hypothetical protein FLJ22031 (FLJ22031), mRNA
NM_025071	Homo sapiens hypothetical protein FLJ21168 (FLJ21168), mRNA
NM_025069	Homo sapiens hypothetical protein FLJ12190 (FLJ12190), mRNA
NM_025067	Homo sapiens hypothetical protein FLJ14299 (FLJ14299) mPN/A
NM_025064	Homo sapiens hypothetical protein FLI14106 (FLI14106) mPNA
NM_025063	Tiomo sapiens hypothetical protein FLJ23604 (FLJ23604) mPNIA
NM 025059	Tiomo sapiens hypothetical protein FLI23550 (FLI23550) mPNIA
	Homo sapiens hypothetical protein FLI23305 (FLI23305) mDNA
	Tiomo sapiens hypothetical protein FLI23189 (FLI23189) mpara
1111_023030	Tromo sapiens hypothetical protein FLI23185 (FLI23185) mDNA
11111_023032	Tiomo sapiens hypothetical protein FL 123074 (FL 123074) mpark
1111_023047	Tromo sapiens hypothetical protein FL 122692 (FL 122692) mpara
1111 025040	Tionio sapiens hypothetical protein FLI22684 (FLI22684) mpara
13/13/1 11/14/13/17/	Homo sapiens hypothetical protein FLJ22595 (FLJ22595), mRNA

NM_025045	Homo sapiens hypothetical protein FLJ22582 (FLJ22582), mRNA
NM_025031	Homo sapiens hypothetical protein FLJ21075 (FLJ21075), mRNA
NM_025030	Homo sapiens hypothetical protein FLJ20972 (FLJ20972), mRNA
NM_025026	Homo sapiens hypothetical protein FLJ14107 (FLJ14107), mRNA
NM_025025	Homo sapiens hypothetical protein FLJ14100 (FLJ14100), mRNA
NM_025024	Homo sapiens hypothetical protein FLJ14082 (FLJ14082), mRNA
NM_025023	Homo sapiens hypothetical protein FLJ14069 (FLJ14069), mRNA
NM_025019	Homo sapiens likely ortholog of mouse tubulin alpha 4 (FLJ13940), mRNA
NM_025012	Homo sapiens hypothetical protein FLJ13769 (FLJ13769), mRNA
NM_025009	Homo sapiens hypothetical protein FLJ13621 (FLJ13621), mRNA
NM_025008	Homo sapiens hypothetical protein FLJ13544 (FLJ13544), mRNA
NM_025006	Homo sapiens hypothetical protein FLJ13373 (FLJ13373), mRNA
NM_025004	Homo sapiens hypothetical protein FLJ13215 (FLJ13215), mRNA
NM_025003	Homo sapiens hypothetical protein FLJ13166 (FLJ13166), mRNA
NM_025002	Homo sapiens hypothetical protein FLJ13162 (FLJ13162), mRNA
NM_025001	Homo sapiens hypothetical protein FLJ13105 (FLJ13105), mRNA
NM_025000	Homo sapiens hypothetical protein FLJ13096 (FLJ13096), mRNA
NM_024997	Homo sapiens hypothetical protein FLJ12668 (FLJ12668), mRNA
NM_024993	Homo sapiens hypothetical protein FLJ12568 (FLJ12568), mRNA
NM_024992	Homo sapiens hypothetical protein FLJ12547 (FLJ12547), mRNA
NM 024989	Homo sapiens hypothetical protein FLJ12377 (FLJ12377), mRNA
NM 024988	Homo sapiens hypothetical protein FLJ12355 (FLJ12355), mRNA
NM 024986	Homo sapiens hypothetical protein FLJ12331 (FLJ12331), mRNA
NM 024980	Homo sapiens hypothetical protein FLJ12132 (FLJ12132), mRNA
NM 024979	Homo sapiens hypothetical protein FLJ12122 (FLJ12122), mRNA
NM 024978	Homo sapiens hypothetical protein FLJ12121 (FLJ12121), mRNA
NM 024971	Homo sapiens hypothetical protein FLJ11726 (FLJ11726), mRNA
NM 024970	Homo sapiens hypothetical protein FLJ11722 (FLJ11722), mRNA
NM 024969	Homo sapiens hypothetical protein FLJ11703 (FLJ11703), mRNA
NM 024966	Homo sapiens hypothetical protein FLJ11598 (FLJ11598), mRNA
NM 024961	Homo sapiens hypothetical protein FLJ11370 (FLJ11370), mRNA
NM 024959	Homo sapiens hypothetical protein FLJ22233 (FLJ22233), mRNA
NM 024957	Homo sapiens hypothetical protein FLJ22686 (FLJ22686), mRNA
NM 024955	Homo sapiens hypothetical protein FLJ23322 (FLJ23322), mRNA
NM 024954	Homo sapiens hypothetical protein FLJ11807 (FLJ11807), mRNA
NM 024952	Homo sapiens hypothetical protein FLJ20950 (FLJ20950), mRNA
NM 024950	Homo sapiens hypothetical protein FLJ12891 (FLJ12891), mRNA
NM 024949	Homo sapiens hypothetical protein FLJ22029 (FLJ22029), mRNA
NM 024948	Homo sapiens hypothetical protein FLJ13397 (FLJ13397), mRNA
NM 024946	Homo sapiens hypothetical protein FLJ21799 (FLJ21799), mRNA
NM 024945	Homo sapiens hypothetical protein FLJ12888 (FLJ12888), mRNA
NM 024943	Homo sapiens hypothetical protein FLJ23235 (FLJ23235), mRNA
NM 024940	Homo sapiens hypothetical protein FLJ21034 (FLJ21034), mRNA
NM 024937	Homo sapiens hypothetical protein FLJ12929 (FLJ12929), mRNA
NM 024936	Homo sapiens hypothetical protein FLJ23024 (FLJ23024), mRNA
NM_024929	Homo sapiens hypothetical protein FLJ23112 (FLJ23112), mRNA
NM 024927	Homo sapiens hypothetical protein FLJ21019 (FLJ21019), mRNA
NM 024926	Homo sapiens hypothetical protein FLJ12571 (FLJ12571), mRNA
NM 024923	Homo sapiens hypothetical protein FLJ22389 (FLJ22389), mRNA
NM 024922	Homo sapiens hypothetical protein FLJ21736 (FLJ21736), mRNA
NM_024921	Homo sapiens hypothetical protein FLJ22792 (FLJ22792), mRNA
L1111 027721	1 Month suprofile hypothetical protein 1 L322172 (1 L322172), hixtyr

377 5 00 10	
NM_024916	
NM_024915	Homo sapiens hypothetical protein FLI13782 (FLI13782) mpnia
NM_024913	Homo sapiens hypothetical protein FLJ21986 (FLJ21986) mPNA
NM_024912	Homo sapiens hypothetical protein FLJ14327 (FLJ14327) mPNA
NM_024910	Tiomo sapiens hypothetical protein FLJ12700 (FLJ12700) mDNA
NM_024902	fromo sapiens hypothetical protein FLI13236 (FLI13236) mDNA
NM_024901	Homo sapiens hypothetical protein FL 122457 (FL 122457) mDNA
NM_024899	Tiomo sapiens hypothetical protein FLI12542 (FLI12542) mPNA
NM_024895	110mo sapiens hypothetical protein FI 123200 (FI 123200) PNIA
NM_024892	From Sapiens hypothetical protein FLI11700 (FLI11700) mDNA
NM_024891	110110 sapiens hypothetical protein FLI11783 (FI I11782) mpN/A
NM_024888	1101110 sapiens hypothetical protein FI I11535 (FI I11535) DAIA
NM_024887	110mo sapiens hypothetical protein FLJ13102 (FLJ13102) mPNA
NM_024884	Homo sapiens hypothetical protein FL 112618 (FL 112619) DATA
NM_024883	Homo sapiens hypothetical protein FLI22202 (FLI22202) mpN/A
NM 024881	110mo sapiens hypothetical protein FL 114251 (FT 114251) DNA
NM_024876	Tiomo sapiens hypothetical protein FLJ12229 (FLJ12229) mPNA
NM 024875	Tiomo sapiens hypothetical protein FL 112921 (FL 112021) DNIA
NM_024872	Homo sapiens hypothetical protein FL 122570 (FI 122570) mDNA
NM_024871	Tiomo sapiens hypothetical protein FLI12748 (FLI12748) mDNA
NM_024869	Homo sapiens hypothetical protein FLI14050 (FLI14050) mDNA
NM_024868	110110 sapiens hypothetical protein FLI14124 (FLI14124) mDNIA
NM_024866 NM_024865	Fiomo sapiens hypothetical protein FLJ21135 (FI I21135) mDNA
NM_024863	FIGURE Sapiens hypothetical protein FLJ12581 (FI J12581) mpN/A
NM_024862	Homo sapiens hypothetical protein FLI21174 (FI I21174) mPNA
NM_024860	Homo sapiens hypothetical protein FLJ13962 (FLJ13962) mDNA
NM_024857	Homo sapiens hypothetical protein FLJ21148 (FLJ21148), mRNA
NM_024855	Homo sapiens hypothetical protein FLJ12735 (FLJ12735), mRNA
NM 024854	Homo sapiens hypothetical protein FLJ12785 (FLJ12785), mRNA
NM_024852	Homo sapiens hypothetical protein FLJ22028 (FLJ22028), mRNA
NM_024850	Homo sapiens hypothetical protein FLJ12765 (FLJ12765), mRNA
NM_024849	Homo sapiens hypothetical protein FLJ21458 (FLJ21458), mRNA Homo sapiens hypothetical protein FLJ21458 (FLJ21458), mRNA
NM_024846	Homo sapiens hypothetical protein FLJ14126 (FLJ14126), mRNA Homo sapiens hypothetical protein FLJ11710 (FLJ11710), mRNA
NM_024845	Homo sapiens hypothetical protein FLJ14154 (FLJ14154), mRNA  Homo sapiens hypothetical protein FLJ14154 (FLJ14154), mRNA
NM_024844	Homo sapiens hypothetical protein FLJ12549 (FLJ12549), mRNA
NM_024843	Homo sapiens duodenal cytochrome b (FLJ23462), mRNA
NM_024838	Homo sapiens hypothetical protein FLJ22002 (FLJ22002), mRNA
NM_024834	Homo sapiens hypothetical protein FLJ13081 (FLJ13081), mRNA
NM_024833	Homo sapiens hypothetical protein FLJ23506 (FLJ23506), mRNA
NM_024830	Homo sapiens hypothetical protein FLJ12443 (FLJ12443), mRNA
NM_024829	Homo sapiens hypothetical protein FLJ22662 (FLJ22662), mRNA Homo sapiens hypothetical protein FLJ22662 (FLJ22662), mRNA
NM_024828	Homo sapiens hypothetical protein FLJ13657 (FLJ13657), mRNA
NM_024827	Homo sapiens hypothetical protein FLJ22237 (FLJ22237), mRNA
NM_024826	110110 sapiens hypothetical protein FLI21159 (FLI21150) DAIA
NM_024825	110mo sapiens hypothetical protein FL 123447 (FI 123447) Data
NM_024824	Homo sapiens hypothetical protein FLJ11806 (FLJ11806), mRNA
NM_024823	Tiomo sapiens hypothetical protein FLJ21596 (FLJ21596) mDNA
NM_024821	from sapiens hypothetical protein FLJ22349 (FLJ22349) mDNA
NM_024818	fromo sapiens hypothetical protein FLI23251 (FI I23251) mDNA
NM_024817	Homo sapiens hypothetical protein FLJ13710 (FLJ13710), mRNA
	(2 2010), IIICIM

NM_024814	Homo sapiens hypothetical protein FLJ23109 (FLJ23109), mRNA
NM_024802	Homo sapiens hypothetical protein FLJ21369 (FLJ21369), mRNA
NM_024801	Homo sapiens hypothetical protein FLJ21551 (FLJ21551), mRNA
NM_024800	Homo sapiens hypothetical protein FLJ23495 (FLJ23495), mRNA
NM_024798	Homo sapiens hypothetical protein FLJ13952 (FLJ13952), mRNA
NM_024794	Homo sapiens hypothetical protein FLJ22408 (FLJ22408), mRNA
NM_024792	Homo sapiens hypothetical protein FLJ22282 (FLJ22282), mRNA
NM_024791	Homo sapiens hypothetical protein FLJ22756 (FLJ22756), mRNA
NM_024790	Homo sapiens hypothetical protein FLJ22490 (FLJ22490), mRNA
NM_024788	Homo sapiens hypothetical protein FLJ21062 (FLJ21062), mRNA
NM_024787	Homo sapiens hypothetical protein FLJ12526 (FLJ12526), mRNA
NM_024786	Homo sapiens hypothetical protein FLJ13153 (FLJ13153), mRNA
NM_024785	Homo sapiens hypothetical protein FLJ22746 (FLJ22746), mRNA
NM_024783	Homo sapiens hypothetical protein FLJ23598 (FLJ23598), mRNA
NM_024782	Homo sapiens hypothetical protein FLJ12610 (FLJ12610), mRNA
NM_024781	Homo sapiens hypothetical protein FLJ23594 (FLJ23594), mRNA
NM_024779	Homo sapiens hypothetical protein FLJ22055 (FLJ22055), mRNA
NM_024778	Homo sapiens hypothetical protein FLJ22612 (FLJ22612), mRNA
NM_024776	Homo sapiens hypothetical protein FLJ21140 (FLJ21140), mRNA
NM_024774	Homo sapiens hypothetical protein FLJ21924 (FLJ21924), mRNA
NM_024770	Homo sapiens hypothetical protein FLJ13984 (FLJ13984), mRNA
NM_024768	Homo sapiens hypothetical protein FLJ12057 (FLJ12057), mRNA
NM_024766	Homo sapiens hypothetical protein FLJ23451 (FLJ23451), mRNA
NM_024765	Homo sapiens hypothetical protein FLJ12401 (FLJ12401), mRNA
NM_024764	Homo sapiens hypothetical protein FLJ14298 (FLJ14298), mRNA
NM_024761	Homo sapiens hypothetical protein FLJ13204 (FLJ13204), mRNA
NM_024759	Homo sapiens hypothetical protein FLJ13955 (FLJ13955), mRNA
NM_024757	Homo sapiens hypothetical protein FLJ12879 (FLJ12879), mRNA
NM_024756	Homo sapiens hypothetical protein FLJ13465 (FLJ13465), mRNA
NM_024755	Homo sapiens hypothetical protein FLJ13213 (FLJ13213), mRNA
NM_024753	Homo sapiens hypothetical protein FLJ11457 (FLJ11457), mRNA
NM_024751	Homo sapiens hypothetical protein FLJ13273 (FLJ13273), mRNA
NM_024748	Homo sapiens hypothetical protein FLJ11539 (FLJ11539), mRNA
NM_024747	Homo sapiens hypothetical protein FLJ22501 (FLJ22501), mRNA
NM_024745	Homo sapiens hypothetical protein FLJ22009 (FLJ22009), mRNA
NM_024743	Homo sapiens hypothetical protein FLJ21934 (FLJ21934), mRNA
NM_024738	Homo sapiens hypothetical protein FLJ21415 (FLJ21415), mRNA
NM_024736	Homo sapiens hypothetical protein FLJ12150 (FLJ12150), mRNA
NM_024735	Homo sapiens hypothetical protein FLJ22477 (FLJ22477), mRNA
NM_024734	Homo sapiens calponin like transmembrane domain protein (calmin), mRNA
NM_024733	Homo sapiens hypothetical protein FLJ14345 (FLJ14345), mRNA
NM_024730	Homo sapiens hypothetical protein FLJ22655 (FLJ22655), mRNA
NM_024729	Homo sapiens hypothetical protein FLJ13881 (FLJ13881), mRNA
NM_024728	Homo sapiens hypothetical protein FLJ11808 (FLJ11808), mRNA
NM_024725	Homo sapiens hypothetical protein FLJ23518 (FLJ23518), mRNA
NM 024724	Homo sapiens hypothetical protein FLJ22332 (FLJ22332), mRNA
NM_024721	Homo sapiens likely ortholog of mouse zinc finger homeodomain 4 (FLJ20980),
_	mRNA
NM_024713	Homo sapiens hypothetical protein FLJ22557 (FLJ22557), mRNA
NM_024712	Homo sapiens engulfment and cell motility 3 (ced-12 homolog, C. elegans)
	(ELMO3), mRNA
NM_024711	Homo sapiens hypothetical protein FLJ22690 (FLJ22690), mRNA

....

۵.

NM_024710	
NM_024710	
NM 024707	Tionio sapiens hypothetical protein El 199551 (El 199551)
NM 024706	110110 Sapiens hypothetical protein FI 113056 (El 112056)
	Tionio sapiens hypothetical profein FI 113470 (ET 112470) DATA
NM_024704	1101110 sapicits hypothetical profess El 122045 (El 122045)
NM_024702	Tiomo sapiens hypothetical profem El 1138/1 (El 1138/1)
NM_024699	Tionio sapiens hypothetical profein El 114007 (El 114007) Data
NM_024697	1101110 Sapiciis flybolnetical profein El 122410 (El 122410)
NM_024696	Tiolio sapielis hypothetical profein El 123059 (El 123059)
NM_024694	Tromo sapiens hypotherical profein El 193191 (CT 199191)
NM_024691	Tionio sapiens hypothetical protein FI 123223 (Er 122222)
NM_024685	Tromo sapiciis hypothetical profein Fl. 123560 (Et. 122560)
NM_024682	Tromo sapicus hypothelical profein ki 113160 oct 113160
NM_024680	1 Tromo sapiciis ilypothelical profein El 173311 (CT 133311)
NM_024679	Tromo sapiens hypothetical profein El 111030 (El 111020) Dazi
NM_024677	Tromo sapiens hypothetical protein El 11/001 (Et 11/001)
NM_024676	Trothe sapicus hypothetical profess El 122020 (Et 122020)
NM_024674	Tromo sapiens hypothetical profess RI 112457 (Et 112457)
NM_024671	Tromo sapiens hypothetical profein El 123/126 (El 122/126)
NM_024669	TIOTHO SAPICHS HYDOLHERICAL profein FI 111705 (TT 111705)
NM_024667	Tromo sapiens hypothetical profein El 112750 (El 112750)
NM_024665	Homo sapiens nuclear receptor co-repressor/HDAC3 complex subunit
NM_024664	Homo sapiens hypothetical protein FI I11838 (FI I11838)
NM_024661	110 The Sapicity Hypothetical protein El 112/126 (ET 112/126)
NM_024660	Tromo sapiens hypothenical profein El 122572 (Et 122572)
NM_024659	Homo sapiens hypothetical protein FLJ11753 (FLJ11753), mRNA Homo sapiens hypothetical protein FLJ11753 (FLJ11753), mRNA
NM_024658	Homo sapiens hypothetical protein FLJ23338 (FLJ23338), mRNA Homo sapiens hypothetical protein FLJ23338 (FLJ23338), mRNA
NM_024657	Tiomo sapiens hypothetical protein El 111565 (El 111565)
NM_024656	Tromo sapiens hypothenical profein El 122220 (Et 122220)
NM_024653	Tromo sapiciis hypothetical profein FL 113002 (ET 112002)
NM_024652	Homo sapiens hypothetical protein FLJ23119 (FLJ23119), mRNA  Homo sapiens hypothetical protein FLJ23119 (FLJ23119), mRNA
NM_024645	Tiomo sapiciis ilypolifetical profess Hi 113942 (Et 113942)
NM_024644	Homo sapiens hypothetical protein FLJ21802 (FLJ21802), mRNA Homo sapiens hypothetical protein FLJ21802 (FLJ21802), mRNA
NM_024643	AXOMO SUPICIS IIVIDII IEI ICAI DEGLES EL 122002 (ET 122002)
NM_024642	Homo sapiens hypothetical protein FLJ21212 (FLJ21212), mRNA  Homo sapiens hypothetical protein FLJ21212 (FLJ21212), mRNA
NM_024639	Homo sapiens hypothetical protein FLJ23393 (FLJ23393), mRNA Homo sapiens hypothetical protein FLJ23393 (FLJ23393), mRNA
NM_024638	1 1101110 Sapicity Hydrights have been fill 110000 (Et 110000)
NM_024635	Tromo sapiens hypothemical profein El 192642 (Et 192642)
NM_024633	Tromo supicità il voluncio di professi el 191976 (Et 191976)
NM_024632	110 mo sapicus hypothenical profess fil 111526 (Et 111526)
NM_024631	azomo sapiciis hyboliiciical profein H1 177742 (Ct 192242)
NM_024630	220110 Suprens hypotherical protein El 170004 (Et toogo ()
NM_024629	Homo sapiens hypothetical protein FLJ23468 (FLJ23468), mRNA Homo sapiens hypothetical protein FLJ23468 (FLJ23468), mRNA
NM_024623	Homo sapiens hypothetical protein FLJ23468 (FLJ23468), mRNA Homo sapiens hypothetical protein FLJ13491 (FLJ13491), mRNA
NM_024620	Homo sapiens hypothetical protein FLJ13491 (FLJ13491), mRNA Homo sapiens hypothetical protein FLJ12586 (FLJ12586), mRNA
NM_024619	Homo sapiens hypothetical protein FLJ12586 (FLJ12586), mRNA Homo sapiens hypothetical protein FLJ12171 (FLJ12171), mRNA
NM_024618	Homo sapiens hypothetical protein FLJ121/1 (FLJ12171), mRNA
NM_024614	Homo sapiens hypothetical protein FLJ21478 (FLJ21478), mRNA Homo sapiens hypothetical protein FLJ13197 (FLJ13197), mRNA Homo sapiens hypothetical protein FLJ13197 (FLJ13197), mRNA
	Homo sapiens hypothetical protein FL 122000 grant Park
NM_024608	Homo sapiens hypothetical protein FLJ22060 (FLJ22060), mRNA Homo sapiens hypothetical protein FLJ22402 (FLJ22402), mRNA
	Pothetical protein FLJ22402 (FLJ22402), mRNA

NM_024607	Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 3B (PPP1R3B), mRNA
NM_024604	Homo sapiens hypothetical protein FLJ21908 (FLJ21908), mRNA
NM 024603	Homo sapiens hypothetical protein FLJ11588 (FLJ11588), mRNA
NM 024599	Homo sapiens hypothetical protein FLJ22341 (FLJ22341), mRNA
NM 024598	Homo sapiens hypothetical protein FLJ13154 (FLJ13154), mRNA
NM 024597	Homo sapiens hypothetical protein FLJ12649 (FLJ12649), mRNA
NM 024596	Homo sapiens hypothetical protein FLJ12847 (FLJ12847), mRNA
NM 024594	Homo sapiens hypothetical protein FLJ12899 (FLJ12899), mRNA
NM 024593	Homo sapiens hypothetical protein FLJ11767 (FLJ11767), mRNA
NM 024592	Homo sapiens hypothetical protein FLJ13352 (FLJ13352), mRNA
NM 024590	Homo sapiens hypothetical protein FLJ23548 (FLJ23548), mRNA
NM 024589	Homo sapiens hypothetical protein FLJ22386 (FLJ22386), mRNA
NM 024588	Homo sapiens hypothetical protein FLJ23584 (FLJ23584), mRNA
NM 024587	Homo sapiens hypothetical protein FLJ22353 (FLJ22353), mRNA
NM 024583	Homo sapiens hypothetical protein FLJ23142 (FLJ23142), mRNA
NM 024582	Homo sapiens hypothetical protein FLJ23056 (FLJ23056), mRNA
NM 024581	Homo sapiens hypothetical protein FLJ13942 (FLJ13942), mRNA
NM 024579	Homo sapiens hypothetical protein FLJ23221 (FLJ23221), mRNA
NM 024578	Homo sapiens hypothetical protein FLJ22709 (FLJ22709), mRNA
NM 024577	Homo sapiens hypothetical protein FLJ13605 (FLJ13605), mRNA
NM 024576	Homo sapiens hypothetical protein FLJ21079 (FLJ21079), mRNA
NM 024575	Homo sapiens hypothetical protein FLJ23467 (FLJ23467), mRNA
NM 024574	Homo sapiens hypothetical protein FLJ23191 (FLJ23191), mRNA
NM 024573	Homo sapiens hypothetical protein FLJ12910 (FLJ12910), mRNA
NM 024572	Homo sapiens hypothetical protein FLJ12691 (FLJ12691), mRNA
NM 024569	Homo sapiens hypothetical protein FLJ21047 (FLJ21047), mRNA
NM 024567	Homo sapiens hypothetical protein FLJ21616 (FLJ21616), mRNA
NM 024564	Homo sapiens hypothetical protein FLJ11715 (FLJ11715), mRNA
NM 024563	Homo sapiens hypothetical protein FLJ14054 (FLJ14054), mRNA
NM 024560	Homo sapiens hypothetical protein FLJ21963 (FLJ21963), mRNA
NM 024558	Homo sapiens hypothetical protein FLJ13920 (FLJ13920), mRNA
NM 024557	Homo sapiens hypothetical protein FLJ11608 (FLJ11608), mRNA
NM 024554	Homo sapiens hypothetical protein FLJ11413 (FLJ11413), mRNA
NM 024548	Homo sapiens hypothetical protein FLJ23047 (FLJ23047), mRNA
NM 024545	Homo sapiens hypothetical protein FLJ12761 (FLJ12761), mRNA
NM 024544	Homo sapiens hypothetical protein FLJ12875 (FLJ12875), mRNA
NM 024541	Homo sapiens hypothetical protein FLJ13114 (FLJ13114), mRNA
NM 024539	Homo sapiens hypothetical protein FLJ23516 (FLJ23516), mRNA
NM 024537	Homo sapiens hypothetical protein FLJ12118 (FLJ12118), mRNA
NM 024536	Homo sapiens hypothetical protein FLJ22678 (FLJ22678), mRNA
NM 024535	Homo sapiens hypothetical protein FLJ22078 (FLJ22078), mRNA
NM 024533	Homo sapiens hypothetical protein FLJ22167 (FLJ22167), mRNA
NM 024533	Homo sapiens hypothetical protein FLJ11856 (FLJ11856), mRNA
NM 024530	Homo sapiens hypothetical protein FLJ23306 (FLJ23306), mRNA
NM 024528	Homo sapiens hypothetical protein FLJ22626 (FLJ22626), mRNA
NM 024527	Homo sapiens hypothetical protein FLJ11743 (FLJ11743), mRNA
NM 024525	Homo sapiens hypothetical protein FLJ22584 (FLJ22584), mRNA
NM 024524	Homo sapiens hypothetical protein FLJ20986 (FLJ20986), mRNA
NM 024521	Homo sapiens hypothetical protein FLJ21459 (FLJ21459), mRNA
NM 024520	Homo sapiens hypothetical protein FLJ21439 (FLJ21439), mRNA  Homo sapiens hypothetical protein FLJ22555 (FLJ22555), mRNA
11111 024320	1 Homo sapiens hypomeneai protein PL322333 (PL322333), filkina

NDV 024510	TT
NM_024519	
NM_024509	Homo sapiens hypothetical protein MGC2656 (MGC2656) mRNA
NM_024506	Homo sapiens hypothetical protein MGC10771 (MGC10771) mRNA
NM_022893	Homo sapiens B-cell CLL/lymphoma 11A (zinc finger protein) (RCI 11A)
ND4 015110	
NM 015113	Homo sapiens KIAA0399 protein (KIAA0399), mRNA
NM_015545	Homo sapiens KIAA0632 protein (KIAA0632) mRNA
NM_020299	Homo sapiens aldo-keto reductase family 1 member B10 (aldose reductase)
NIM 002208	(MACIDIO), IIRNA
NM 003308	Homo sapiens testis specific protein, Y-linked (TSPY), mRNA
NM_024339	Tiomo sapiens hypothetical protein MGC2655 (MGC2655) mPNA
NM 024334 NM 024328	Homo sapiens hypothetical protein MGC3222 (MGC3222) mPNA
	Homo sapiens hypothetical protein MGC2652 (MGC2652) mPNA
NM_024327 NM_024323	Homo sapiens hypothetical protein MGC2508 (MGC2508), mPNA
	Homo sapiens hypothetical protein MGC11271 (MGC11271) mPNA
NM_024322 NM_024320	Homo sapiens hypothetical protein MGC11266 (MGC11266), mDNA
NM 024320 NM 024319	Homo sapiens hypothetical protein MGC11242 (MGC11242) mRNA
NM_024314	Homo sapiens hypothetical protein MGC4174 (MGC4174) mPNA
NM_024314 NM_024313	Homo sapiens hypothetical protein MGC4294 (MGC4294) mPNIA
	Homo sapiens hypothetical protein MGC3731 (MGC3731) mPNIA
NM_024310 NM_024303	Homo sapiens hypothetical protein MGC4090 (MGC4090) mDNA
NM_024297	Fiorio sapiens hypothetical protein MGC4161 (MGC4161) mDNA
	1 Homo sapiens hypothetical protein MGC2941 (MGC2941) mPNIA
NM 024293	Homo sapiens hypothetical protein MGC3035 (MGC3035) mPNIA
NM_023003	Homo sapiens transmembrane 6 superfamily member 1 (TM6CE1) DNA
NM_015254	Homo sapielis kinesin family member 13B (KIF13B) mRNA
NM_015127 NM_024033	Homo sapiens Mid-1-related chloride channel 1 (KIA A0761) mDNA
NM 024122	Homo sapiens hypothetical protein MGC5242 (MGC5242) mpNA
NM_024122	Homo sapiens hypothetical protein MGC4825 (MGC4825) mPNA
NM_024121	Homo sapiens hypothetical protein FLJ20979 (FLJ20979) mPNA
NM_024119	Homo sapiens hypothetical protein FLI11354 (FLI11354) mpNIA
NM_024117	Homo sapiens hypothetical protein MGC2745 (MGC2745) mDNA
NM_024111	Homo sapiens hypothetical protein MGC4309 (MGC/4300) DNIA
NM_024111	Hollo sapiens hypothetical protein MGC4504 (MGC4504) mPNIA
NM_024108	Homo sapiens hypothetical protein MGC2654 (MGC2654) mDNA
NM_024108	Homo sapiens hypothetical protein MGC2650 (MGC2650) DNIA
NM_024107	Homo sapiens hypothetical protein MGC3123 (MGC3123) mDNA
NM_024104	1101110 sapiens hypothetical protein MGC2663 (MGC2663) mDNIA
NM_024104 NM_024102	Homo sapiens hypothetical protein MGC2747 (MGC2747) DNIA
NM_024097	Homo sapiens hypothetical protein MGC2722 (MGC2722) DNA
NM_024094	110110 sapiciis hypothetical profein MGC055 (MCC055) DATA
NM_024093	Homo sapiens hypothetical protein MGC5528 (MGC5528) mDNA
NM_024090	Tionio sapiens hypothetical profein MGC5500 (MGC5500) DNA
NM_024086	Homo sapiens hypothetical protein MGC5487 (I CF) mPNA
NM_024085	Homo sapiens hypothetical protein MGC3329 (MGC3329) mDNIA
NM_024080	fromo sapiens hypothetical protein FLI22169 (FLI22169) mpNA
NM_024080	nomo sapiens hypothetical protein MGC2849 (MGC2849) mDNIA
NM_024076	nomo sapiens hypothetical protein MGC2628 (MGC2628) mpNA
	Tiomo sapiens hypothetical protein MGC3169 (MGC3169) mpNIA
NM_024071 NM_024070	nomo sapiens hypothetical protein MGC2550 (MGC2550) mpNi
	Homo sapiens hypothetical protein MGC2463 (MGC2463) mDNA
1111 024009	Homo sapiens hypothetical protein MGC2749 (MGC2749), mRNA

NM_024068	Homo sapiens hypothetical protein MGC2731 (MGC2731), mRNA
NM_024065	Homo sapiens hypothetical protein MGC3062 (MGC3062), mRNA
NM_024061	Homo sapiens hypothetical protein MGC5521 (MGC5521), mRNA
NM_024058	Homo sapiens hypothetical protein MGC5590 (MGC5590), mRNA
NM 024057	Homo sapiens hypothetical protein MGC5585 (MGC5585), mRNA
NM 024053	Homo sapiens hypothetical protein MGC861 (MGC861), mRNA
NM 024050	Homo sapiens hypothetical protein MGC2594 (MGC2594), mRNA
NM 024049	Homo sapiens hypothetical protein MGC5566 (MGC5566), mRNA
NM 024048	Homo sapiens hypothetical protein MGC3020 (MGC3020), mRNA
NM 024046	Homo sapiens hypothetical protein MGC8407 (MGC8407), mRNA
NM 024045	Homo sapiens nucleolar protein GU2 (GU2), mRNA
NM 024041	Homo sapiens hypothetical protein MGC3180 (MGC3180), mRNA
NM 024039	Homo sapiens hypothetical protein MGC2488 (MGC2488), mRNA
NM 024038	Homo sapiens hypothetical protein MGC2803 (MGC2803), mRNA
NM 024037	Homo sapiens hypothetical protein MGC2603 (MGC2603), mRNA
NM 024032	Homo sapiens hypothetical protein MGC3130 (MGC3130), mRNA
NM 024031	Homo sapiens hypothetical protein MGC3121 (MGC3121), mRNA
NM 024028	Homo sapiens hypothetical protein MGC3265 (MGC3265), mRNA
NM 024027	Homo sapiens hypothetical protein MGC3279 similar to collectins (MGC3279),
TVIVI_024027	mRNA
NM 024025	Homo sapiens hypothetical protein MGC1136 (MGC1136), mRNA
NM 024006	Homo sapiens hypothetical protein IMAGE3455200 (IMAGE3455200), mRNA
NM 015653	Homo sapiens DKFZP566F0546 protein (DKFZP566F0546), mRNA
NM 015147	Homo sapiens KIAA0582 protein (KIAA0582), mRNA
NM 016481	Homo sapiens hypothetical protein (HSPC219), mRNA
NM 023940	Homo sapiens hypothetical protein MGC2827 (MGC2827), mRNA
NM 023938	Homo sapiens hypothetical protein MGC2742 (MGC2742), mRNA
NM 023931	Homo sapiens hypothetical protein MGC2474 (MGC2474), mRNA
NM_015517	Homo sapiens MBD2 (methyl-CpG-binding protein)-interacting zinc finger
) D 6 015540	protein (MIZF), mRNA
NM_015540	Homo sapiens DKFZP727M111 protein (DKFZP727M111), mRNA
NM_015043	Homo sapiens KIAA0676 protein (KIAA0676), mRNA
NM_023934	Homo sapiens hypothetical protein MGC2495 (MGC2495), mRNA
NM_023928	Homo sapiens hypothetical protein FLJ12389 similar to acetoacetyl-CoA synthetase (FLJ12389), mRNA
NM 023926	Homo sapiens hypothetical protein FLJ12895 (FLJ12895), mRNA
NM 023924	Homo sapiens hypothetical protein FLJ13441 (FLJ13441), mRNA
NM 020239	Homo sapiens small protein effector 1 of Cdc42 (SPEC1), mRNA
NM_012069	Homo sapiens ATPase, (Na+)/K+ transporting, beta 4 polypeptide (ATP1B4),
NM 023112	mRNA Homo sapiens hypothetical protein FLJ21916 (FLJ21916), mRNA
NM 015324	Homo sapiens KIAA0409 protein (KIAA0409), mRNA
NM 023079	Homo sapiens hypothetical protein FLJ13855 (FLJ13855), mRNA
NM 023077	Homo sapiens hypothetical protein FLJ12439 (FLJ12439), mRNA
NM 023075	Homo sapiens hypothetical protein FLJ11585 (FLJ11585), mRNA
NM 023074	Homo sapiens hypothetical protein FLJ12644 (FLJ12644), mRNA
NM 023073	Homo sapiens hypothetical protein FLJ13231 (FLJ13231), mRNA
NM_023071	Homo sapiens hypothetical protein FLJ13117 (FLJ13117), mRNA
NM 012319	Homo sapiens LIV-1 protein, estrogen regulated (LIV-1), mRNA
NM_023012	Homo sapiens LIV-1 protein, estrogen regulated (LIV-1), fireNA  Homo sapiens hypothetical protein FLJ11021 similar to splicing factor,
NWI_023012	arginine/serine-rich 4 (FLJ11021), mRNA
NM_023008	Homo sapiens hypothetical protein FLJ12949 (FLJ12949), mRNA

NM_023007	
NM_022918	Homo sapiens hypothetical protein FLI22104 (FLI22104) mpht
NM_022914	Homo sapiens hypothetical protein 24432 (24432) mRNA
NM_022912	Homo sapiens hypothetical protein FLJ13110 (FLJ13110), mRNA
NM_022907	Homo sapiens hypothetical protein FLJ23053 (FLJ23053), mRNA
NM_022905	Homo sapiens hypothetical protein FLJ12572 (FLJ12572), mRNA
NM 022901	Homo saniens hypothetical protein FL31202 (FL312572), mRNA
NM_022898	Homo sapiens hypothetical protein FLJ21302 (FLJ21302), mRNA  Homo sapiens R-cell CLI (hymphory 11R (i.e., 5)
	Homo sapiens B-cell CLL/lymphoma 11B (zinc finger protein) (BCL11B), mRNA
NM_022841	Homo sapiens hypothetical protein FLJ12994 (FLJ12994), mRNA
NM 022840	Homo sapiens hypothetical protein FLJ23017 (FLJ23017), mRNA  Homo sepiens hypothetical protein FLJ23017 (FLJ23017), mRNA
NM 022834	Homo saniens hypothetical protein FL 122017 (FL 122017), mRNA
NM_022832	Homo sapiens hypothetical protein FLJ22215 (FLJ22215), mRNA
NM_022827	Homo sapiens hypothetical protein FLJ12552 (FLJ12552), mRNA Homo sapiens hypothetical
NM 022826	Homo sapiens hypothetical protein FLJ21347 (FLJ21347), mRNA
NM_022823	Homo sapiens axotrophin (AXOT), mRNA  Homo sapiens hypothetical actions and the sapiens hypothetical actions are sapiens as a sapiens and the sapiens are sapiens as a sapiens are sapiens are sapiens as a sapiens are sapiens ar
NM_022781	Homo sapiens hypothetical protein FLJ22362 (FLJ22362), mRNA
NM 022780	Homo sapiens hypothetical protein FLJ21343 (FLJ21343), mRNA
NM_022788	Tionio sapiciis hypothetical profess ki 113010 (Et 113010)
1111_022110	Homo sapiens hypothetical protein DKFZp434L0117 (DKFZP434L0117), mRNA
NM_022777	11111111
NM_022771	Homo sapiens hypothetical protein FLJ14117 (FLJ14117), mRNA
NM_022770	Tiomo sapiens hypothetical protein FLI12085 (FI 112085) mDNIA
NM_022769	Tromo sapiens hypothetical protein FL 113912 (FL 113012) DNIA
NM_022767	Tromo sapiens hypothetical protein FL 121868 (FL 121868) mDNA
NM_022766	Fromo Sapiens hypothetical protein FL 112484 (FT 112484)
NM 022763	From Sapiens hypothetical protein FL 123239 (FT 123239) mpNIA
NM_022762	Tionio sapiens hypothetical profein FI 123300 (FI 123300) DAIA
NM_022759	1 Hollo Sapiens hypothetical protein FI 122318 (FI 122318) DAIA
NM_022754	Tiomo sapiens hypothetical protein FLI21865 (FLI21865) mDNA
NM_022752	fromo sapiens hypothetical profem FLI12876 (FLI112876) DNIA
	Homo sapiens hypothetical protein FI 122050 (FI 122050) DATA
NM_022751	110mo sapiens hypothetical protein FLI21610 (FLI21610) -DNIA
NM_022750	Tromo sapiens hypothetical protein FL 122693 (FL 122603) mDNIA
NM_022747	Homo sapiens hypothetical protein FI 122559 (FI 122559)
NM_022744	Hollo sapiens hypothetical protein FI 113868 (FI 113969) DNA
NM_022743	TIOMO Sapiens hypothetical profem El 121080 (El 121080) Para
NM_022741	Tiomo sapiens hypothetical protein FLI11850 (FLI11850) DNA
NM_022736	Tiomo sapiens hypothetical protein FI II/153 (FI II/152) DAI/
NM_022734	Tiomo sapiens hypothetical profein FI 120850 (FI 120850) DAIA
NM_022731	Tromo sapiens similar to rat nuclear ubjouitous casein kingse 2 OHICKS
ND COORSE	1 1111/11/11
NM_022727	Homo sapiens Hpall tiny fragments locus 9C (HTF9C), mRNA
NM_012197	110th Sapiens 1400 GIPase activating protein (GAP and contraction)
NDA 015126	(CIL CENTY), IIII(IA
NM_015136	Homo sapiens KIAA0246 protein (stab1), mRNA
NM_022659	Homo sapiens likely ortholog of mouse early Breell factor 2 (EL 111500)
NM_022571	residence of the support of the supp
ND4 001001	(HOMINELLEO), MKNA
NM_021024	Homo sapiens high-mobility group (nonhistone chromosomal) protein 17-like 1
ND C CCCC	(ALIGITALI), IIICIVA
NM_019884	Homo sapiens glycogen synthase kinase 3 alpha (GSK3A), mRNA
NM_021034	Homo sapiens interferon induced transmembrane protein 3 (1-8U) (IFITM3),
	protein 5 (1-80) (IFIIM3),

	mRNA
NM 022445	Homo sapiens thiamin pyrophosphokinase 1 (TPK1), mRNA
NM 022495	Homo sapiens hypothetical protein FLJ12799 (FLJ12799), mRNA
NM 022494	Homo sapiens hypothetical protein FLJ21952 (FLJ21952), mRNA
NM 022492	Homo sapiens hypothetical protein FLJ12788 (FLJ12788), mRNA
NM 022488	Homo sapiens PC3-96 protein (PC3-96), mRNA
NM 022480	Homo sapiens hypothetical protein FLJ12587 (FLJ12587), mRNA
NM 022474	Homo sapiens hypothetical protein FLJ12615 similar to membrane protein,
	palmitoylated 3 (MAGUK p55 subfamily member 5) (FLJ12615), mRNA
NM 022455	Homo sapiens androgen receptor-associated coregulator 267 (ARA267), mRNA
NM 022452	Homo sapiens hypothetical protein FLJ11618 (FLJ11618), mRNA
NM 022448	Homo sapiens hypothetical protein FLJ21817 similar to Rhoip2 (FLJ21817),
_	mRNA
NM 022373	Homo sapiens hypothetical protein FLJ22313 (FLJ22313), mRNA
NM 022370	Homo sapiens hypothetical protein FLJ21044 similar to Rbig1 (FLJ21044),
_	mRNA
NM_022368	Homo sapiens praja 1 (PJA1), mRNA
NM_022366	Homo sapiens hypothetical protein FLJ23182 (FLJ23182), mRNA
NM_022361	Homo sapiens popeye protein 3 (POP3), mRNA
NM_022360	Homo sapiens human epididymis-specific 3 beta (HE3-BETA), mRNA
NM_022342	Homo sapiens kinesin family member 9 (KIF9), mRNA
NM_022372	Homo sapiens G protein beta subunit-like (GBL), mRNA
NM_022158	Homo sapiens fructosamine-3-kinase (FN3K), mRNA
NM_022137	Homo sapiens secreted modular calcium-binding protein 1 (SMOC1), mRNA
NM_022118	Homo sapiens cutaneous T-cell lymphoma tumor antigen se70-2 (SE70-2),
	mRNA
NM_022116	Homo sapiens fidgetin-like 1 (FIGNL1), mRNA
NM_022103	Homo sapiens hypothetical zinc finger protein FLJ14011 (FLJ14011), mRNA
NM_022070	Homo sapiens hypothetical protein FLJ22087 (FLJ22087), mRNA
NM_022065	Homo sapiens hypothetical protein FLJ21877 (FLJ21877), mRNA
NM_021970	Homo sapiens mitogen-activated protein kinase kinase 1 interacting protein 1
77.5 04.0004	(MAP2K1IP1), mRNA
NM_019081	Homo sapiens KIAA0430 gene product (KIAA0430), mRNA
NM_021981	Homo sapiens pre-T/NK cell associated protein (1D12A), mRNA
NM_020121	Homo sapiens UDP-glucose ceramide glucosyltransferase-like 2 (UGCGL2), mRNA
NM_006683	Homo sapiens human epididymis-specific 3 alpha (HE3-ALPHA), mRNA
NM 006077	Homo sapiens calcium binding atopy-related autoantigen 1 (CBARA1), mRNA
NM 021934	Homo sapiens hypothetical protein FLJ11773 (FLJ11773), mRNA
NM 021933	Homo sapiens hypothetical protein FLJ12438 (FLJ12438), mRNA
NM 021930	Homo sapiens Rad50-interacting protein 1 (FLJ11785), mRNA
NM 021929	Homo sapiens hypothetical protein FLJ21613 similar to rat corneal wound
	healing related protein (FLJ21613), mRNA
NM 007272	Homo sapiens chymotrypsin C (caldecrin) (CTRC), mRNA
NM_004237	Homo sapiens thyroid hormone receptor interactor 13 (TRIP13), mRNA
NM_003849	Homo sapiens succinate-CoA ligase, GDP-forming, alpha subunit (SUCLG1),
	mRNA
NM_021648	Homo sapiens KIAA0721 protein (KIAA0721), mRNA
NM_021831	Homo sapiens hypothetical protein FLJ21839 (FLJ21839), mRNA
NM_021827	Homo sapiens hypothetical protein FLJ23514 (FLJ23514), mRNA
NM_021195	Homo sapiens claudin 6 (CLDN6), mRNA
NM_018947	Homo sapiens cytochrome c (HCS), mRNA
	(/)

NM_021732	Homo sapiens hypothetical protein PP5395 (PP5395), mRNA
NM_021730	Homo sapiens hypothetical protein PP1044 (PP1044), mRNA
NM_021643	Homo sapiens GS3955 protein (GS3955), mRNA
NM_015180	Homo sapiens synaptic nuclei expressed gene 2 (SYNE-2), mRNA
NM_021633	Homo sapiens kelch-like protein C3IP1 (C3IP1), mRNA
NM_021629	Homo sapiens guanine nucleotide binding protein beta subunit 4 (GNB4), mRNA
NM_021627	Homo sapiens sentrin-specific protease (SENP2), mRNA
NM_021626	Homo sapiens likely homolog of rat and mouse retinoid-inducible serine carboxypeptidase (RISC), mRNA
NM_021622	Homo sapiens pleckstrin homology domain-containing, family A (phosphoinositide binding specific) member 1 (PLEKHA1), mRNA
NM_012408	Homo sapiens protein kinase C binding protein 1 (PRKCBP1), mRNA
NM 021252	Homo sapiens RAB18, member RAS oncogene family (RAB18), mRNA
NM 020806	Homo sapiens gephyrin (GPHN), mRNA
NM 021258	Homo sapiens interleukin 22 receptor (IL22R), mRNA
NM_021235	Homo sapiens epidermal growth factor receptor substrate EPS15R (EPS15R), mRNA
NM_021204	Homo sapiens E-1 enzyme (MASA), mRNA
NM_021191	Homo sapiens neurogenic differentiation 4 (NEUROD4), mRNA
NM_021178	Homo sapiens enhancer of invasion 10 (HEI10), mRNA
NM_021127	Homo sapiens phorbol-12-myristate-13-acetate-induced protein 1 (PMAIP1), mRNA
NM_021114	Homo sapiens serine protease inhibitor, Kazal type, 2 (acrosin-trypsin inhibitor) (SPINK2), mRNA
NM_021103	Homo sapiens thymosin, beta 10 (TMSB10), mRNA
NM_006435	Homo sapiens interferon induced transmembrane protein 2 (1-8D) (IFITM2), mRNA
NM_021073	Homo sapiens bone morphogenetic protein 5 (BMP5), mRNA
NM_003142	Homo sapiens Sjogren syndrome antigen B (autoantigen La) (SSB), mRNA
NM_003888	Homo sapiens aldehyde dehydrogenase 1 family, member A2 (ALDH1A2), mRNA
NM_013234	Homo sapiens muscle specific gene (M9), mRNA
NM_021067	Homo sapiens KIAA0186 gene product (KIAA0186), mRNA
NM_021020	Homo sapiens leucine zipper, putative tumor suppressor 1 (LZTS1), mRNA
NM_021025	Homo sapiens homeo box 11-like 2 (HOX11L2), mRNA
NM_021003	Homo sapiens protein phosphatase 1A (formerly 2C), magnesium-dependent, alpha isoform (PPM1A), mRNA
NM_020674	Homo sapiens cytochrome P450 monooxygenase (CYP-M), mRNA
NM_019612	Homo sapiens hypothetical protein R30953_1 (R30953_1), mRNA
NM_020904	Homo sapiens pleckstrin homology domain-containing, family A (phosphoinositide binding specific) member 4 (PLEKHA4), mRNA
NM_020686	Homo sapiens NPD009 protein (NPD009), mRNA
NM 020684	Homo sapiens NPD007 protein (NPD007), mRNA
NM_020683	Homo sapiens AD026 protein (AD026), mRNA
NM_020679	Homo sapiens AD023 protein (AD023), mRNA
NM_020677	Homo sapiens HSCARG protein (HSCARG), mRNA
NM 020675	Homo sapiens AD024 protein (AD024) mRNA
NM_020675 NM_020673	Homo sapiens AD024 protein (AD024), mRNA
NM_020673	Homo sapiens AD024 protein (AD024), mRNA Homo sapiens RAB22A, member RAS oncogene family (RAB22A), mRNA
NM_020673 NM_020660	Homo sapiens AD024 protein (AD024), mRNA Homo sapiens RAB22A, member RAS oncogene family (RAB22A), mRNA Homo sapiens connexin-36 (CX36), mRNA
NM_020673	Homo sapiens AD024 protein (AD024), mRNA Homo sapiens RAB22A, member RAS oncogene family (RAB22A), mRNA

NB ( 019424	YI coniona coliett
NM_018434	Homo sapiens goliath protein (GP), mRNA
NM_020437	Homo sapiens similar to aspartate beta hydroxylase (ASPH) (LOC57168),
37.5 000504	mRNA
NM_020524	Homo sapiens hematopoietic PBX-interacting protein (HPIP), mRNA
NM_018638	Homo sapiens ethanolamine kinase (EKI1), mRNA
NM_016326	Homo sapiens chemokine-like factor 1 (CKLF1), mRNA
NM_016951	Homo sapiens chemokine-like factor 1 (CKLF1), mRNA
NM_020143	Homo sapiens putatative 28 kDa protein (LOC56902), mRNA
NM_020141	Homo sapiens protein x 013 (AD-020), mRNA
NM_020122	Homo sapiens potassium channel modulatory factor (PCMF), mRNA
NM_018843	Homo sapiens mitochondrial carrier family protein (MCFP), mRNA
NM 018840	Homo sapiens putative Rab5-interacting protein (RIP5), mRNA
NM 016303	Homo sapiens pp21 homolog (LOC51186), mRNA
NM 016300	Homo sapiens cyclic AMP-regulated phosphoprotein, 21 kD (ARPP-21), mRNA
NM 016299	Homo sapiens likely ortholog of mouse heat shock protein, 70 kDa 4
_	(LOC51182), mRNA
NM 013259	Homo sapiens neuronal protein (NP25), mRNA
NM_005064	Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 23
	(SCYA23), mRNA
NM 013260	Homo sapiens transcriptional regulator protein (HCNGP), mRNA
NM 020433	Homo sapiens hypothetical protein LOC57158 (LOC57158), mRNA
NM 020410	Homo sapiens CGI-152 protein (CGI-152), mRNA
NM 020401	Homo sapiens nuclear pore complex protein (NUP107), mRNA
NM 020400	Homo sapiens G protein-coupled receptor 92 (GPR92), mRNA
NM 020397	Homo sapiens CamKI-like protein kinase (LOC57118), mRNA
NM 020388	Homo sapiens CATX-15 protein (CATX-15), mRNA
NM 020386	Homo sapiens HRAS-like suppressor (HRASLS), mRNA
NM 020361	Homo sapiens carboxypeptidase B precursor (CPAH), mRNA
NM 020357	Homo sapiens PEST-containing nuclear protein (pcnp), mRNA
NM 020345	Homo sapiens I-kappa-B-interacting Ras-like protein 1 (KBRAS1), mRNA
NM 020360	Homo sapiens phospholipid scramblase 3 (PLSCR3), mRNA
NM 020348	
	Homo sapiens cyclin M1 (CNNM1), mRNA
NM_000888	Homo sapiens integrin, beta 6 (ITGB6), mRNA
NM_020181	Homo sapiens myelin proteolipid protein-like protein (PLPL), mRNA
NM_020144	Homo sapiens poly(A) polymerase beta (testis specific) (PAPOLB), mRNA
NM_020202	Homo sapiens Nit protein 2 (NIT2), mRNA
NM_020250	Homo sapiens MOST2 protein (MOST2), mRNA
NM_020237	Homo sapiens MOST-1 protein (MOST-1), mRNA
NM_020234	Homo sapiens x 009 protein (MDS009), mRNA
NM_020128	Homo sapiens nuclear protein double minute 1 (MDM1), mRNA
NM_020169	Homo sapiens latexin protein (LXN), mRNA
NM_020133	
	mRNA
NM_020241	Homo sapiens sema domain, transmembrane domain (TM), and cytoplasmic
	domain, (semaphorin) 6B (SEMA6B), mRNA
NM_020163	Homo sapiens semaphorin sem2 (LOC56920), mRNA
NM_020199	Homo sapiens HTGN29 protein (HTGN29), mRNA
NM_020197	Homo sapiens HSKM-B protein (HSKM-B), mRNA
NM_020200	Homo sapiens HHGP protein (HHGP), mRNA
NM_020195	Homo sapiens HCDI protein (HCDI), mRNA
NM 020198	Homo sapiens GK001 protein (GK001), mRNA
NM_020133 NM_020241 NM_020163 NM_020199 NM_020197 NM_020200 NM_020195	Homo sapiens lysophosphatidic acid acyltransferase-delta (LPAAT-delta), mRNA  Homo sapiens sema domain, transmembrane domain (TM), and cytoplasmic domain, (semaphorin) 6B (SEMA6B), mRNA  Homo sapiens semaphorin sem2 (LOC56920), mRNA  Homo sapiens HTGN29 protein (HTGN29), mRNA  Homo sapiens HSKM-B protein (HSKM-B), mRNA  Homo sapiens HHGP protein (HHGP), mRNA  Homo sapiens HCDI protein (HCDI), mRNA

NM_020119	Homo sapiens hypothetical protein FLB6421 (FLB6421), mRNA
NM_020162	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 33 (DDX33),
	mRNA
NM_020215	Homo sapiens hypothetical protein DKFZp761F2014 (DKFZp761F2014).
	mRNA
NM_020221	Homo sapiens hypothetical protein DKFZp547I224 (DKFZp547I224), mRNA
NM_020217	Homo sapiens hypothetical protein DKFZp547I014 (DKFZp547I014), mRNA
NM_020161	Homo sapiens hypothetical protein DKFZp547H025 (DKFZp547H025), mRNA
NM 020186	Homo sapiens DC11 protein (DC11), mRNA
NM 020205	Homo sapiens cellular zinc finger anti-NF-kappaB Cezanne (CEZANNE),
_	mRNA
NM 019887	Homo sapiens second mitochondria-derived activator of caspase (SMAC),
_	mRNA
NM 019892	Homo sapiens phosphatidylinositol (4,5) bisphosphate 5-phosphatase homolog;
	phosphatidylinositol polyphosphate 5-phosphatase type IV (PPI5PIV), mRNA
NM 019885	Homo sapiens cytochrome P450 retinoid metabolizing protein (P450RAI-2),
	mRNA
NM_019845	Homo sapiens candidate mediator of the p53-dependent G2 arrest (REPRIMO),
	mRNA
NM 019853	Homo sapiens protein phosphatase 4 regulatory subunit 2 (PPP4R2), mRNA
NM_013301	Homo sapiens protein predicted by clone 23882 (HSU79303), mRNA
NM 013300	Homo sapiens protein predicted by clone 23733 (HSU79274), mRNA
NM 013296	Homo sapiens LGN protein (HSU54999), mRNA
NM 013293	
NM 013310	Homo sapiens transformer-2 alpha (htra-2 alpha) (HSU53209), mRNA
NM 018975	Homo sapiens hypothetical protein (AF038169), mRNA
NM 019082	Homo sapiens TRF2-interacting telomeric RAP1 protein (RAP1), mRNA
NM 019020	Homo sapiens putative nucleolar RNA helicase (NOH61), mRNA Homo sapiens hypothetical protein (FLJ20748), mRNA
NM 019058	Homo sapiens HIF-1 responsive RTP801 (FLJ20500), mRNA
NM 019056	Homo sapiens neuronal protein 17.3 (P17.3), mRNA
NM_019042	
NM 019061	Homo sapiens hypothetical protein (FLJ20485), mRNA
14141_019001	Homo sapiens phosphatidylinositol-3 phosphate 3-phosphatase adaptor subunit (3-PAP), mRNA
NM 018986	
NM_019034	Homo sapiens hypothetical protein (FLJ20356), mRNA
14141_017034	Homo sapiens ras homolog gene family, member F (in filopodia) (ARHF), mRNA
NM_019062	Homo sapiens hypothetical protein (FLJ20225), mRNA
NM 019038	Homo sapiens hypothetical protein (FLJ1045), mRNA  Homo sapiens hypothetical protein (FLJ11045), mRNA
NM_019044	
NM 018180	Homo sapiens hypothetical protein (FLJ10996), mRNA
14147_010100	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 32 (DDX32), mRNA
NM 019028	
1,11,1 0,1,20,70	Homo sapiens hypothetical protein similar to ankyrin repeat-containing priotein
NM 019014	AKR1 (FLJ10852), mRNA
14147 013014	Homo sapiens similar to DNA-directed RNA polymerase I (135 kDa) (Rpo1-2),
NM 019023	mRNA
NM_018162	Homo sapiens hypothetical protein (FLJ10640), mRNA
NM_018162 NM_019067	Homo sapiens hypothetical protein FLJ10633 (FLJ10633), mRNA
	Homo sapiens hypothetical protein (FLJ10613), mRNA
NM_019057	Homo sapiens hypothetical protein (FLJ10404), mRNA
NM_018846	Homo sapiens SBBI26 protein (SBBI26), mRNA
NM_016483	Homo sapiens hypothetical protein (HSPC226), mRNA
NM_018400	Homo sapiens voltage-gated sodium channel beta-3 subunit (scn3b gene)

	(US A 2/2206) mDN/A
NM 018700	(HSA243396), mRNA
NM 018547	Homo sapiens tripartite motif-containing 36 (TRIM36), mRNA
NM 018546	Homo sapiens hypothetical protein PRO2964 (PRO2964), mRNA
NM 018544	Homo sapiens hypothetical protein PRO2958 (PRO2958), mRNA
	Homo sapiens hypothetical protein PRO2949 (PRO2949), mRNA
NM_018634	Homo sapiens hypothetical protein PRO2893 (PRO2893), mRNA
NM_018543	Homo sapiens hypothetical protein PRO2859 (PRO2859), mRNA
NM_018542	Homo sapiens hypothetical protein PRO2834 (PRO2834), mRNA
NM_018538	Homo sapiens erythroblast membrane-associated protein (ERMAP), mRNA
NM_018534	Homo sapiens hypothetical protein PRO2714 (PRO2714), mRNA
NM_018530	Homo sapiens hypothetical protein PRO2521 (PRO2521), mRNA
NM_018627	Homo sapiens hypothetical protein PRO2405 (PRO2405), mRNA
NM_018523	Homo sapiens hypothetical protein PRO2325 (PRO2325), mRNA
NM_018519	Homo sapiens hypothetical protein PRO2266 (PRO2266), mRNA
NM_018517	Homo sapiens hypothetical protein PRO2214 (PRO2214), mRNA
NM_018621	Homo sapiens hypothetical protein PRO2198 (PRO2198), mRNA
NM_018619	Homo sapiens hypothetical protein PRO2133 (PRO2133), mRNA
NM_018618	Homo sapiens hypothetical protein PRO2121 (PRO2121), mRNA
NM_018616	Homo sapiens hypothetical protein PRO2037 (PRO2037), mRNA
NM_018512	Homo sapiens hypothetical protein PRO2015 (PRO2015), mRNA
NM_018610	Homo sapiens hypothetical protein PRO1942 (PRO1942), mRNA
NM_018510	Homo sapiens hypothetical protein PRO1866 (PRO1866), mRNA
NM_018507	Homo sapiens hypothetical protein PRO1843 (PRO1843), mRNA
NM_018606	Homo sapiens hypothetical protein PRO1787 (PRO1787), mRNA
NM_018589	Homo sapiens hypothetical protein PRO1635 (PRO1635), mRNA
NM_018587	Homo sapiens hypothetical protein PRO1617 (PRO1617), mRNA
NM_018503	Homo sapiens hypothetical protein PRO1598 (PRO1598), mRNA
NM_018586	Homo sapiens hypothetical protein PRO1584 (PRO1584), mRNA
NM_018502	Homo sapiens hypothetical protein PRO1580 (PRO1580), mRNA
NM_018603	Homo sapiens hypothetical protein PRO1496 (PRO1496), mRNA
NM_018584	Homo sapiens hypothetical protein PRO1489 (PRO1489), mRNA
NM_018582	Homo sapiens hypothetical protein PRO1483 (PRO1483), mRNA
NM_018602	Homo sapiens DnaJ (Hsp40) homolog, subfamily A, member 4 (DNAJA4),
	mRNA
NM_018578	Homo sapiens hypothetical protein PRO1257 (PRO1257), mRNA
NM_018576	Homo sapiens hypothetical protein PRO1163 (PRO1163), mRNA
NM_018497	Homo sapiens hypothetical protein PRO1048 (PRO1048), mRNA
NM_018565	Homo sapiens hypothetical protein PRO0899 (PRO0899), mRNA
NM_018562	Homo sapiens hypothetical protein PRO0386 (PRO0386), mRNA
NM_018590	Homo sapiens hypothetical protein PRO0082 (PRO0082), mRNA
NM_018667	Homo sapiens sphingomyelin phosphodiesterase 3, neutral membrane (neutral
	sphingomyelinase II) (SMPD3), mRNA
NM_017544	Homo sapiens transcription factor NRF (NRF), mRNA
NM_018468	Homo sapiens uncharacterized hematopoietic stem/progenitor cells protein
	MDS033 (MDS033), mRNA
NM_018467	Homo sapiens uncharacterized hematopoietic stem/progenitor cells protein
	MDS032 (MDS032), mRNA
NM_018464	Homo sapiens uncharacterized hematopoietic stem/progenitor cells protein
	MDS029 (MDS029), mRNA
NM_018688	Homo sapiens bridging integrator 3 (BIN3), mRNA
NM_018686	Homo sapiens CMP-N-acetylneuraminic acid synthase (CMAS), mRNA
NM_018446	Homo sapiens glycosyltransferase AD-017 (AD-017), mRNA

NM_018416	Homo sapiens FOXJ2 forkhead factor (FHX), mRNA
NM_018407	Homo sapiens putative integral membrane transporter (LC27), mRNA
NM_018472	Homo sapiens uncharacterized hypothalamus protein HT011 (HT011), mRNA
NM_018471	Homo sapiens uncharacterized hypothalamus protein HT010 (HT010), mRNA
NM_018470	Homo sapiens uncharacterized hypothalamus protein HT009 (HT009), mRNA
NM_018469	Homo sapiens uncharacterized hypothalamus protein HT008 (HT008), mRNA
NM_017523	Homo sapiens XIAP associated factor-1 (HSXIAPAF1), mRNA
NM_017514	Homo sapiens SEX gene (HSSEXGENE), mRNA
NM_017512	Homo sapiens rTS beta protein (HSRTSBETA), mRNA
NM_016536	Homo sapiens HSPC059 protein (HSPC059), mRNA
NM_018553	Homo sapiens ELG protein (HSA277841), mRNA
NM_018403	Homo sapiens transcription factor (SMIF gene) (HSA275986), mRNA
NM_018404	Homo sapiens centaurin, alpha 2 (CENTA2), mRNA
NM_018401	Homo sapiens gene for serine/threonine protein kinase (HSA250839), mRNA
NM_017582	Homo sapiens NICE-5 protein (HSA243666), mRNA
NM 018684	Homo sapiens hepatocellular carcinoma-associated antigen 127 (HCA127),
	mRNA
NM 018477	Homo sapiens uncharacterized hypothalamus protein HARP11 (HARP11).
	mRNA
NM_018652	Homo sapiens golgin-like protein (GLP), mRNA
NM_017962	Homo sapiens hypothetical protein FLJ20825 (FLJ20825), mRNA
NM_017961	Homo sapiens hypothetical protein FLJ20813 (FLJ20813), mRNA
NM_017960	Homo sapiens hypothetical protein FLJ20808 (FLJ20808), mRNA
NM_017959	Homo sapiens hypothetical protein FLJ20802 (FLJ20802), mRNA
NM_017958	Homo sapiens hypothetical protein FLJ20783 (FLJ20783), mRNA
NM_017957	Homo sapiens epsin 3 (FLJ20778), mRNA
NM_017956	Homo sapiens hypothetical protein FLJ20772 (FLJ20772), mRNA
NM_017950	Homo sapiens hypothetical protein FLJ20753 (FLJ20753), mRNA
NM_017949	Homo sapiens hypothetical protein FLJ20739 (FLJ20739), mRNA
NM_017946	Homo sapiens hypothetical protein FLJ20731 (FLJ20731), mRNA
NM_017953	Homo sapiens hypothetical protein FLJ20729 (FLJ20729), mRNA
NM_017943	Homo sapiens hypothetical protein FLJ20725 (FLJ20725), mRNA
NM_017941	Homo sapiens hypothetical protein FLJ20721 (FLJ20721), mRNA
NM_017938	Homo sapiens hypothetical protein FLJ20716 (FLJ20716), mRNA
NM_017937	Homo sapiens hypothetical protein FLJ20712 (FLJ20712), mRNA
NM_017932	Homo sapiens hypothetical protein FLJ20700 (FLJ20700), mRNA
NM_017929	Homo sapiens hypothetical protein FLJ20695 (FLJ20695), mRNA
NM_017928	Homo sapiens hypothetical protein FLJ20694 (FLJ20694), mRNA
NM_017925	Homo sapiens hypothetical protein FLJ20686 (FLJ20686), mRNA
NM_017920	Homo sapiens hypothetical protein FLJ20654 (FLJ20654), mRNA
NM_017919	Homo sapiens hypothetical protein FLJ20651 (FLJ20651), mRNA
NM_017918	Homo sapiens hypothetical protein FLJ20647 (FLJ20647), mRNA
NM_017917	Homo sapiens hypothetical protein FLJ20644 (FLJ20644), mRNA
NM_017916	Homo sapiens hypothetical protein FLJ20643 (FLJ20643), mRNA
NM_017915	Homo sapiens hypothetical protein FLJ20641 (FLJ20641), mRNA
NM_017912	Homo sapiens hypothetical protein FLJ20637 (FLJ20637), mRNA
NM_017909	Homo sapiens hypothetical protein FLJ20627 (FLJ20627), mRNA
NM_017907	Homo sapiens hypothetical protein FLJ20625 (FLJ20625), mRNA
NM_017903	Homo sapiens hypothetical protein FLJ20618 (FLJ20618), mRNA
NM_017901	Homo sapiens two-pore channel 1, homolog (KIAA1169), mRNA
NM_017900	Homo sapiens hypothetical protein FLJ20608 (FLJ20608), mRNA
NM_017899	Homo sapiens hypothetical protein FLJ20607 (TSC), mRNA

	<del></del>
NM_017897	Homo sapiens hypothetical protein FLJ20604 (FLJ20604), mRNA
NM_017894	Homo sapiens hypothetical protein FLJ20595 (FLJ20595), mRNA
NM_017893	Homo sapiens sema domain, immunoglobulin domain (Ig), transmembrane
	domain (TM) and short cytoplasmic domain, (semaphorin) 4G (SEMA4G),
3 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2	mRNA
NM_017891	Homo sapiens hypothetical protein FLJ20584 (FLJ20584), mRNA
NM_017885	Homo sapiens hypothetical protein FLJ20568 (FLJ20568), mRNA
NM_017881	Homo sapiens hypothetical protein FLJ20559 (FLJ20559), mRNA
NM_017876	Homo sapiens hypothetical protein FLJ20552 (FLJ20552), mRNA
NM_017873	Homo sapiens hypothetical protein FLJ20548 (FLJ20548), mRNA
NM_017868	Homo sapiens hypothetical protein FLJ20535 (FLJ20535), mRNA
NM_017866	Homo sapiens hypothetical protein FLJ20533 (FLJ20533), mRNA
NM_017863	Homo sapiens hypothetical protein FLJ20527 (FLJ20527), mRNA
NM_017860	Homo sapiens hypothetical protein FLJ20519 (FLJ20519), mRNA
NM_017858	Homo sapiens hypothetical protein FLJ20516 (FLJ20516), mRNA
NM_017856	Homo sapiens hypothetical protein FLJ20514 (FLJ20514), mRNA
NM_017854	Homo sapiens hypothetical protein FLJ20512 (FLJ20512), mRNA
NM_017853	Homo sapiens hypothetical protein FLJ20511 (FLJ20511), mRNA
NM_017851	Homo sapiens hypothetical protein FLJ20509 (FLJ20509), mRNA
NM_017848	Homo sapiens hypothetical protein FLJ20506 (FLJ20506), mRNA
NM_017843	Homo sapiens breast carcinoma amplified sequence 4 (BCAS4), mRNA
NM_017836	Homo sapiens hypothetical protein FLJ20473 (FLJ20473), mRNA
NM_017834	Homo sapiens hypothetical protein FLJ20464 (FLJ20464), mRNA
NM_017831	Homo sapiens hypothetical protein FLJ20456 (FLJ20456), mRNA
NM_017828	Homo sapiens hypothetical protein FLJ20452 (FLJ20452), mRNA
NM_017825	Homo sapiens hypothetical protein FLJ20446 (FLJ20446), mRNA
NM_017824	Homo sapiens hypothetical protein FLJ20445 (FLJ20445), mRNA
NM_017819	Homo sapiens hypothetical protein FLJ20432 (FLJ20432), mRNA
NM_017817	Homo sapiens hypothetical protein FLJ20429 (FLJ20429), mRNA
NM_017816	Homo sapiens hypothetical protein FLJ20425 (FLJ20425), mRNA
NM_017814	Homo sapiens hypothetical protein FLJ20422 (FLJ20422), mRNA
NM_017813	Homo sapiens hypothetical protein FLJ20421 (FLJ20421), mRNA
NM_017812	Homo sapiens hypothetical protein FLJ20420 (FLJ20420), mRNA
NM_017808	Homo sapiens hypothetical protein FLJ20413 (FLJ20413), mRNA
NM_017805	Homo sapiens hypothetical protein FLJ20401 (FLJ20401), mRNA
NM_017803	Homo sapiens hypothetical protein FLJ20399 (FLJ20399), mRNA
NM_017801 NM_017799	Homo sapiens hypothetical protein FLJ20396 (FLJ20396), mRNA
NM 017793	Homo sapiens hypothetical protein FLJ20392 (FLJ20392), mRNA
NM 017791	Homo sapiens hypothetical protein FLJ20374 (FLJ20374), mRNA
NM_017787	Homo sapiens hypothetical protein FLJ20371 (FLJ20371), mRNA
NM 017782	Homo sapiens hypothetical protein FLJ20154 (FLJ20154), mRNA
NM_017781	Homo sapiens hypothetical protein FLJ20360 (FLJ20360), mRNA Homo sapiens hypothetical protein FLJ20359 (FLJ20359), mRNA
NM_017779	Homo sapiens hypothetical protein FLJ20359 (FLJ20359), mRNA  Homo sapiens hypothetical protein FLJ20354 (FLJ20354), mRNA
NM 017777	
NM 017776	Homo sapiens hypothetical protein FLJ20345 (FLJ20345), mRNA
NM 017773	Homo sapiens hypothetical protein FLJ20344 (FLJ20344), mRNA
NM 017769	Homo sapiens hypothetical protein FLJ20340 (FLJ20340), mRNA
NM 017767	Homo sapiens hypothetical protein FLJ20333 (FLJ20333), mRNA
NM 017766	Homo sapiens hypothetical protein FLJ20327 (FLJ20327), mRNA
NM 017765	Homo sapiens hypothetical protein FLJ20321 (FLJ20321), mRNA
NM 017763	Homo sapiens hypothetical protein FLJ20320 (FLJ20320), mRNA Homo sapiens hypothetical protein FLJ20315 (FLJ20315), mRNA
T414T 01 1 102	Latomo sapiens hypotheticai protein FLJ20313 (FLJ20313), MKNA

NM_017761 Homo sapiens hypothetical protein FLJ20312 (FLJ20312), mRNA NM_017760 Homo sapiens hypothetical protein FLJ20311 (FLJ20311), mRNA NM_017755 Homo sapiens hypothetical protein FLJ20303 (FLJ20303), mRNA NM_017752 Homo sapiens hypothetical protein FLJ20298 (FLJ20298), mRNA NM_017750 Homo sapiens hypothetical protein FLJ20296 (FLJ20296), mRNA NM_017746 Homo sapiens hypothetical protein FLJ20287 (FLJ20287), mRNA NM_017745 Homo sapiens hypothetical protein FLJ20285 (FLJ20285), mRNA NM_017742 Homo sapiens hypothetical protein FLJ20281 (FLJ20281), mRNA
NM 017755 Homo sapiens hypothetical protein FLJ20303 (FLJ20303), mRNA NM 017752 Homo sapiens hypothetical protein FLJ20298 (FLJ20298), mRNA NM 017750 Homo sapiens hypothetical protein FLJ20296 (FLJ20296), mRNA NM 017746 Homo sapiens hypothetical protein FLJ20287 (FLJ20287), mRNA NM 017745 Homo sapiens hypothetical protein FLJ20285 (FLJ20285), mRNA NM 017742 Homo sapiens hypothetical protein FLJ20281 (FLJ20281), mRNA
NM 017752 Homo sapiens hypothetical protein FLJ20298 (FLJ20298), mRNA NM 017750 Homo sapiens hypothetical protein FLJ20296 (FLJ20296), mRNA NM 017746 Homo sapiens hypothetical protein FLJ20287 (FLJ20287), mRNA NM 017745 Homo sapiens hypothetical protein FLJ20285 (FLJ20285), mRNA NM 017742 Homo sapiens hypothetical protein FLJ20281 (FLJ20281), mRNA
NM 017750 Homo sapiens hypothetical protein FLJ20296 (FLJ20296), mRNA NM 017746 Homo sapiens hypothetical protein FLJ20287 (FLJ20287), mRNA NM 017745 Homo sapiens hypothetical protein FLJ20285 (FLJ20285), mRNA NM 017742 Homo sapiens hypothetical protein FLJ20281 (FLJ20281), mRNA
NM017746Homo sapiens hypothetical protein FLJ20287 (FLJ20287), mRNANM017745Homo sapiens hypothetical protein FLJ20285 (FLJ20285), mRNANM017742Homo sapiens hypothetical protein FLJ20281 (FLJ20281), mRNA
NM_017745 Homo sapiens hypothetical protein FLJ20285 (FLJ20285), mRNA NM_017742 Homo sapiens hypothetical protein FLJ20281 (FLJ20281), mRNA
NM_017742 Homo sapiens hypothetical protein FLJ20281 (FLJ20281), mRNA
NM_017741 Homo sapiens hypothetical protein FLJ20280 (FLJ20280), mRNA
NM_017739 Homo sapiens O-linked mannose beta1,2-N-acetylglucosaminyltransferase
(FLJ20277), mRNA  NM 017737 Homo sapiens hypothetical protein FLJ20275 (FLJ20275) mRNA
NM 017729 Homo sapiens hypothetical protein FLJ20258 (FLJ20258), mRNA
NM_017728 Homo sapiens hypothetical protein FLJ20255 (FLJ20255), mRNA
NM_017727 Homo sapiens hypothetical protein FLJ20254 (FLJ20254), mRNA
NM_017724 Homo sapiens leucine rich repeat (in FLII) interacting protein 2 (LRRFIP2), mRNA
NM_017721 Homo sapiens hypothetical protein FLJ20241 (FLJ20241), mRNA
NM_017713 Homo sapiens hypothetical protein FLJ20211 (FLJ20211), mRNA
NM_017712 Homo sapiens hypothetical protein FLJ20208 (FLJ20208), mRNA
NM_017710 Homo sapiens hypothetical protein FLJ20203 (FLJ20203), mRNA
NM 017708 Homo sapiens hypothetical protein FLJ20200 (FLJ20200), mRNA
NM_017707 Homo sapiens hypothetical protein FLJ20199 (FLJ20199), mRNA
NM_017706 Homo sapiens hypothetical protein FLJ20195 (FLJ20195), mRNA
NM_017705 Homo sapiens hypothetical protein FLJ20190 (FLJ20190), mRNA
NM_017703 Homo sapiens hypothetical protein FLJ20188 (FLJ20188), mRNA
NM_017702 Homo sapiens hypothetical protein FLJ20186 (FLJ20186), mRNA
NM_017700 Homo sapiens hypothetical protein FLJ20184 (FLJ20184), mRNA
NM_017696 Homo sapiens hypothetical protein FLJ20170 (FLJ20170), mRNA
NM_017694 Homo sapiens hypothetical protein FLJ20160 (FLJ20160), mRNA
NM_017693 Homo sapiens hypothetical protein FLJ20159 (FLJ20159), mRNA
NM_017691 Homo sapiens hypothetical protein FLJ20156 (FLJ20156), mRNA
NM_017689 Homo sapiens hypothetical protein FLJ20151 (FLJ20151), mRNA
NM_017688 Homo sapiens hypothetical protein FLJ20150 (FLJ20150), mRNA
NM_017685 Homo sapiens hypothetical protein FLJ20139 (FLJ20139), mRNA
NM_017684 Homo sapiens hypothetical protein FLJ20136 (FLJ20136), mRNA
NM_017682 Homo sapiens hypothetical protein FLJ20132 (FLJ20132), mRNA
NM_017681 Homo sapiens hypothetical protein FLJ20130 (FLJ20130), mRNA
NM_017679 Homo sapiens hypothetical protein FLJ20128 (FLJ20128), mRNA
NM_017674 Homo sapiens hypothetical protein FLJ20123 (FLJ20123), mRNA
NM_017664 Homo sapiens hypothetical protein FLJ20093 (FLJ20093), mRNA
NM_017661 Homo sapiens hypothetical protein FLJ20086 (FLJ20086), mRNA
NM_017660 Homo sapiens hypothetical protein FLJ20085 (FLJ20085), mRNA
NM_017658   Homo sapiens hypothetical protein FLJ20081 (FLJ20081), mRNA
NM_017656 Homo sapiens hypothetical protein FLJ20079 (FLJ20079), mRNA
NM_017655 Homo sapiens hypothetical protein FLJ20075 (FLJ20075), mRNA
NM_017654 Homo sapiens hypothetical protein FLJ20073 (FLJ20073), mRNA
NM 017653 Homo sapiens hypothetical protein FLJ20071 (FLJ20071), mRNA
NM_017651 Homo sapiens hypothetical protein FLJ20069 (FLJ20069), mRNA
NM_017650 Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 9A
(PPP1R9A), mRNA
NM_017649 Homo sapiens cyclin M2 (CNNM2), mRNA

NM_017644	Homo sapiens hypothetical protein FLJ20059 (FLJ20059), mRNA
NM 017643	Homo sapiens hypothetical protein FLJ20055 (FLJ20055), mRNA
NM 017639	Homo sapiens hypothetical protein FLJ20047 (FLJ20047), mRNA
NM 017638	Homo sapiens hypothetical protein FLJ20045 (FLJ20045), mRNA
NM 017633	Homo sapiens hypothetical protein FLJ20037 (FLJ20037), mRNA
NM 017631	Homo sapiens hypothetical protein FLJ20035 (FLJ20035), mRNA
NM 017630	Homo sapiens hypothetical protein FLJ20034 (FLJ20034), mRNA
NM 017627	Homo sapiens hypothetical protein FLJ20030 (FLJ20030), mRNA
NM 017626	Homo sapiens DnaJ (Hsp40) homolog, subfamily B, member 12 (DNAJB12),
	mRNA
NM 017621	Homo sapiens hypothetical protein FLJ20013 (FLJ20013), mRNA
NM 017618	Homo sapiens hypothetical protein FLJ20006 (FLJ20006), mRNA
NM 017617	Homo sapiens hypothetical protein FLJ20005 (FLJ20005), mRNA
NM 017615	Homo sapiens hypothetical protein FLJ20003 (FLJ20003), mRNA
NM 018394	Homo sapiens hypothetical protein FLJ11342 (FLJ11342), mRNA
NM 018393	Homo sapiens hypothetical protein FLJ11336 (FLJ11336), mRNA
NM 018391	Homo sapiens hypothetical protein FLJ11328 (FLJ11328), mRNA
NM 018389	Homo sapiens GDP-fucose transporter 1 (FLJ11320), mRNA
NM 018388	Homo sapiens hypothetical protein FLJ11316 (FLJ11316), mRNA
NM 018386	Homo sapiens hypothetical protein FLJ11305 (FLJ11305), mRNA
NM 018383	Homo sapiens hypothetical protein FLJ11294 (FLJ11294), mRNA
NM 018380	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 28 (DDX28),
1.1.1_010500	mRNA
NM 018379	Homo sapiens hypothetical protein FLJ11280 (FLJ11280), mRNA
NM 018376	Homo sapiens hypothetical protein FLJ11275 (FLJ11275), mRNA
NM 018375	Homo sapiens hypothetical protein FLJ11274 (FLJ11274), mRNA
NM 018374	Homo sapiens hypothetical protein FLJ11273 (FLJ11273), mRNA
NM 018372	Homo sapiens hypothetical protein FLJ11269 (FLJ11269), mRNA
NM 018370	Homo sapiens hypothetical protein FLJ11259 (FLJ11259), mRNA
NM 018366	Homo sapiens hypothetical protein FLJ11230 (FLJ11230), mRNA
NM 018365	Homo sapiens hypothetical protein FLJ11222 (FLJ11222), mRNA
NM_018360	Homo sapiens hypothetical protein FLJ11209 (FLJ11209), mRNA
NM 018359	Homo sapiens hypothetical protein FLJ11200 (FLJ11200), mRNA
NM 018357	Homo sapiens hypothetical protein FLJ11196 (FLJ11196), mRNA
NM 018356	Homo sapiens hypothetical protein FLJ11193 (FLJ11193), mRNA
NM 018355	Homo sapiens hypothetical protein FLJ11191 (FLJ11191), mRNA
NM 018351	Homo sapiens hypothetical protein FLJ11183 (FLJ11183), mRNA
NM 018350	Homo sapiens hypothetical protein FLJ11181 (FLJ11181), mRNA
NM 018349	Homo sapiens hypothetical protein FLJ11175 (FLJ11175), mRNA
NM 018348	Homo sapiens hypothetical protein FLJ11171 (FLJ11171), mRNA
NM 018346	Homo sapiens hypothetical protein FLJ11164 (FLJ11164), mRNA
NM_018344	Homo sapiens hypothetical protein FLJ11160 (FLJ11160), mRNA
NM 018343	Homo sapiens hypothetical protein FLJ11159 (FLJ11159), mRNA
NM 018342	Homo sapiens hypothetical protein FLJ11155 (FLJ11155), mRNA
NM 018338	Homo sapiens hypothetical protein FLJ11142 (FLJ11142), mRNA
NM 018335	Homo sapiens hypothetical protein FLJ11132 (FLJ11132), mRNA
NM 018329	Homo sapiens hypothetical protein FLJ11117 (FLJ11117), mRNA
NM 018328	Homo sapiens hypothetical protein FLJ11113 (FLJ11113), mRNA
NM 018326	Homo sapiens hypothetical protein FLJ11110 (FLJ11110), mRNA
NM_018324	Homo sapiens hypothetical protein FLJ11106 (FLJ11106), mRNA
NM_018323	Homo sapiens hypothetical protein FLJ11105 (FLJ11105), mRNA
NM 018321	Homo sapiens hypothetical protein FLJ11100 (FLJ11100), mRNA
	1 - Acomo capicilla il producti a protein i La i i i i i i (i La i i i i i i), i intiga

375 2522	
NM_018316	Homo sapiens hypothetical protein FLJ11078 (FLJ11078), mRNA
NM_018314	Homo sapiens hypothetical protein FLJ11068 (FLJ11068), mRNA
NM_018309	Homo sapiens hypothetical protein FLJ11046 (FLJ11046), mRNA
NM_018308	Homo sapiens hypothetical protein FLJ11042 (FLJ11042), mRNA
NM_018307	Homo sapiens hypothetical protein FLJ11040 (FLJ11040), mRNA
NM_018306	Homo sapiens hypothetical protein FLJ11036 (FLJ11036), mRNA
NM_018304	Homo sapiens hypothetical protein FLJ11029 (FLJ11029), mRNA
NM_018302	Homo sapiens hypothetical protein FLJ11017 (FLJ11017), mRNA
NM_018299	Homo sapiens hypothetical protein FLJ11011 (FLJ11011), mRNA
NM_018297	Homo sapiens peptide:N-glycanase similar to yeast PNG1 (FLJ11005), mRNA
NM_018296	Homo sapiens hypothetical protein FLJ11004 (FLJ11004), mRNA
NM_018294	Homo sapiens hypothetical protein FLJ10998 (FLJ10998), mRNA
NM_018292	Homo sapiens hypothetical protein FLJ10989 (FLJ10989), mRNA
NM_018289	Homo sapiens hypothetical protein FLJ10979 (FLJ10979), mRNA
NM_018288	Homo sapiens hypothetical protein FLJ10975 (FLJ10975), mRNA
NM_018279	Homo sapiens hypothetical protein FLJ10936 (FLJ10936), mRNA
NM_018275	Homo sapiens hypothetical protein FLJ10925 (FLJ10925), mRNA
NM_018271	Homo sapiens hypothetical protein FLJ10916 (FLJ10916), mRNA
NM_018264	Homo sapiens hypothetical protein FLJ10900 (FLJ10900), mRNA
NM_018261	Homo sapiens Sec3-like (SEC3), mRNA
NM_018260	Homo sapiens hypothetical protein FLJ10891 (FLJ10891), mRNA
NM_018259	Homo sapiens hypothetical protein FLJ10890 (FLJ10890), mRNA
NM_018250	Homo sapiens hypothetical protein FLJ10871 (FLJ10871), mRNA
NM_018248	Homo sapiens hypothetical protein FLJ10858 (FLJ10858), mRNA
NM_018247 NM_018246	Homo sapiens hypothetical protein FLJ10856 (FLJ10856), mRNA
NM 018243	Homo sapiens hypothetical protein FLJ10853 (FLJ10853), mRNA
NM 018238	Homo sapiens hypothetical protein FLJ10849 (FLJ10849), mRNA
NM 018235	Homo sapiens hypothetical protein FLJ10842 (FLJ10842), mRNA
NM 018234	Homo sapiens hypothetical protein FLJ10830 (FLJ10830), mRNA
NM_018231	Homo sapiens hypothetical protein FLJ10829 (FLJ10829), mRNA
NM 018229	Homo sapiens hypothetical protein FLJ10815 (FLJ10815), mRNA Homo sapiens hypothetical protein FLJ10813 (FLJ10813), mRNA
NM_018228	Homo sapiens hypothetical protein FLJ10813 (FLJ10813), mRNA Homo sapiens hypothetical protein FLJ10811 (FLJ10811), mRNA
NM_018227	Homo sapiens hypothetical protein FLJ10801 (FLJ10811), mRNA  Homo sapiens hypothetical protein FLJ10808 (FLJ10808), mRNA
NM 018224	Homo sapiens hypothetical protein FLJ10803 (FLJ10803), mRNA  Homo sapiens hypothetical protein FLJ10803 (FLJ10803), mRNA
NM_018222	Homo sapiens parvin, alpha (PARVA), mRNA
NM_018221	Homo sapiens chromosome 2 open reading frame 6 (C2orf6), mRNA
NM_018216	Homo sapiens hypothetical protein FLJ10782 (FLJ10782), mRNA
NM_018215	Homo sapiens hypothetical protein FLJ10782 (FLJ10782), mRNA
NM_018214	Homo sapiens LAP (leucine-rich repeats and PDZ) and no PDZ protein (LANO),
	mRNA
NM_018210	Homo sapiens hypothetical protein FLJ10769 (FLJ10769), mRNA
NM_018208	Homo sapiens hypothetical protein FLJ10761 (FLJ10761), mRNA
NM_018203	Homo sapiens hypothetical protein FLJ10748 (FLJ10748), mRNA
NM_018201	Homo sapiens hypothetical protein FLI10743 (FLI10743) mRNA
NM_018199	Homo sapiens hypothetical protein FLJ10738 (FLJ10738), mRNA
NM_018198	Homo sapiens hypothetical protein FLJ10737 (FLJ10737), mRNA
NM_018196	Homo sapiens epsilon-trimethyllysine hydroxylase (FLJ10727) mRNA
NM_018195	Homo sapiens hypothetical protein FLJ10726 (FLJ10726), mRNA
NM_018190	Homo sapiens hypothetical protein FLJ10715 (FLJ10715), mRNA
NM_018189	Homo sapiens hypothetical protein FLJ10713 (FLJ10713), mRNA

NM_018183	Homo sapiens hypothetical protein FLJ10701 (FLJ10701), mRNA
NM_018182	Homo sapiens hypothetical protein FLJ10700 (FLJ10700), mRNA
NM_018181	Homo sapiens hypothetical protein FLJ10697 (FLJ10697), mRNA
NM_018176	Homo sapiens hypothetical protein FLJ10675 (FLJ10675), mRNA
NM_018174	Homo sapiens chromosome 19 open reading frame 5 (C19orf5), mRNA
NM_018173	Homo sapiens hypothetical protein FLJ10665 (FLJ10665), mRNA
NM_018172	Homo sapiens hypothetical protein FLJ10661 (FLJ10661), mRNA
NM_018170	Homo sapiens hypothetical protein FLJ10656 (FLJ10656), mRNA
NM_018168	Homo sapiens hypothetical protein FLJ10650 (FLJ10650), mRNA
NM_018167	Homo sapiens hypothetical protein FLJ10648 (FLJ10648), mRNA
NM_018166	Homo sapiens hypothetical protein FLJ10647 (FLJ10647), mRNA
NM_018163	Homo sapiens hypothetical protein FLJ10634 (FLJ10634), mRNA
NM 018157	Homo sapiens hypothetical protein FLJ10620 (FLJ10620), mRNA
NM_018156	Homo sapiens hypothetical protein FLJ10619 (FLJ10619), mRNA
NM 018155	Homo sapiens hypothetical protein FLJ10618 (FLJ10618), mRNA
NM 018154	Homo sapiens hypothetical protein FLJ10604 (FLJ10604), mRNA
NM 018150	Homo sapiens hypothetical protein FLJ10597 (FLJ10597), mRNA
NM 018149	Homo sapiens hypothetical protein FLJ10587 (FLJ10587), mRNA
NM 018148	Homo sapiens hypothetical protein FLJ10583 (FLJ10583), mRNA
NM 018146	Homo sapiens hypothetical protein FLJ10581 (FLJ10581), mRNA
NM 018145	Homo sapiens hypothetical protein FLJ10579 (FLJ10579), mRNA
NM 018143	Homo sapiens hypothetical protein FLJ10572 (FLJ10572), mRNA
NM 018140	Homo sapiens hypothetical protein FLJ10565 (FLJ10565), mRNA
NM 018139	Homo sapiens hypothetical protein FLJ10563 (FLJ10563), mRNA
NM 018138	Homo sapiens hypothetical protein FLJ10560 (FLJ10560), mRNA
NM 018132	Homo sapiens hypothetical protein FLJ10545 (FLJ10545), mRNA
NM 018130	Homo sapiens hypothetical protein FLJ10539 (FLJ10539), mRNA
NM 018129	Homo sapiens hypothetical protein FLJ10535 (FLJ10535), mRNA
NM 018128	Homo sapiens hypothetical protein FLJ10534 (FLJ10534), mRNA
NM 018126	Homo sapiens hypothetical protein FLJ10525 (FLJ10525), mRNA
NM_018125	Homo sapiens hypothetical protein FLJ10521 (FLJ10521), mRNA
NM_018121	Homo sapiens hypothetical protein FLJ10512 (FLJ10512), mRNA
NM_018118	Homo sapiens hypothetical protein FLJ10508 (FLJ10508), mRNA
NM 018115	Homo sapiens hypothetical protein FLJ10498 (FLJ10498), mRNA
NM 018113	Homo sapiens lipocalin-interacting membrane receptor (LIMR), mRNA
NM_018111	Homo sapiens hypothetical protein FLJ10490 (FLJ10490), mRNA
NM_018110	Homo sapiens hypothetical protein FLJ10488 (FLJ10488), mRNA
NM_018109	Homo sapiens hypothetical protein FLJ10486 (FLJ10486), mRNA
NM_018108	Homo sapiens hypothetical protein FLJ10483 (FLJ10483), mRNA
NM_018105	Homo sapiens hypothetical protein FLJ10477 (FLJ10477), mRNA
NM_018104	Homo sapiens hypothetical protein FLJ10474 (FLJ10474), mRNA
NM_018096	Homo sapiens hypothetical protein similar to beta-transducin family (FLJ10458),
	mRNA
NM_018095	Homo sapiens hypothetical protein FLJ10450 (FLJ10450), mRNA
NM_018089	Homo sapiens hypothetical protein FLJ10415 (FLJ10415), mRNA
NM_018088	Homo sapiens hypothetical protein FLJ10408 (FLJ10408), mRNA
NM_018084	Homo sapiens hypothetical protein FLJ10392 (FLJ10392), mRNA
NM_018083	Homo sapiens zinc finger protein 358 (ZNF358), mRNA
NM_018082	Homo sapiens hypothetical protein FLJ10388 (FLJ10388), mRNA
NM_018081	Homo sapiens hypothetical protein FLJ10385 (FLJ10385), mRNA
NM 018080	Homo sapiens hypothetical protein FLJ10381 (FLJ10381), mRNA
<del></del>	Myponicular protein 120 10001 (120 10001), HILLIA

NM_018077	TT
NM_018071	Homo sapiens hypothetical protein FLJ10377 (FLJ10377), mRNA
NM 018071	Homo sapiens hypothetical protein FLJ10357 (FLJ10357) mPNA
14141_010000	Homo sapiens likely ortholog of mouse piwi like homolog 1 (Drosophila)-like
NM 018067	1 (P.E. 10331), INKNA
NM_018066	Homo sapiens hypothetical protein FLJ10350 (FLJ10350), mRNA
NM 018065	Homo sapiens hypothetical protein FLJ10349 (FLJ10349), mRNA
NM 018061	Homo sapiens hypothetical protein FLJ10346 (FLJ10346), mRNA
NM_018056	Homo sapiens hypothetical protein FLJ10330 (FLJ10330), mRNA
NM 018055	Homo sapiens hypothetical protein FLJ10315 (FLJ10315) mRNA
NM_018048	Homo sapiens hypothetical protein FLJ10314 (FLJ10314), mRNA
NM_018045	Homo sapiens hypothetical protein FLJ10292 (FLJ10292), mRNA
NM_018042	Homo sapiens hypothetical protein FLJ10276 (FLJ10276), mRNA
NM 018037	Homo sapiens hypothetical protein FLJ10260 (FLJ10260), mRNA
NM_018036	Homo sapiens hypothetical protein FLJ10244 (FLJ10244), mRNA
NM_018029	Homo sapiens hypothetical protein FLJ10242 (FLJ10242), mRNA
NM_018027	Homo sapiens hypothetical protein FLJ10213 (FLJ10213), mRNA
NM_018024	Homo sapiens hypothetical protein FLJ10210 (FLJ10210), mRNA
NM 018022	Homo sapiens hypothetical protein FLJ10204 (FLJ10204), mRNA
NM 018022	Homo sapiens hypothetical protein FLJ10199 (FLJ10199), mRNA
NM_018014	Homo sapiens hypothetical protein FLJ10188 (FLJ10188), mRNA
14141_016014	Homo sapiens B-cell CLL/lymphoma 11A (zinc finger protein) (BCL11A), mRNA
NM 018013	mxnA
NM_018012	Homo sapiens hypothetical protein FLJ10159 (FLJ10159), mRNA
NM_018005	Homo sapiens hypothetical protein FLJ10157 (FLJ10157), mRNA
NM_017998	Homo sapiens hypothetical protein FLJ10139 (FLJ10139), mRNA
NM_017996	Homo sapiens hypothetical protein FLJ10110 (FLJ10110), mRNA
NM_017986	Homo sapiens hypothetical protein FLJ10103 (FLJ10103), mRNA
NM 017985	Homo sapiens hypothetical protein FLJ10060 (FLJ10060), mRNA
NM_017984	Homo sapiens hypothetical protein FLJ10058 (FLJ10058), mRNA
NM_017983	Homo sapiens hypothetical protein FLJ10057 (FLJ10057), mRNA
NM_017982	Homo sapiens hypothetical protein FLJ10055 (FLJ10055), mRNA
NM_017980	Homo sapiens hypothetical protein FLJ10052 (FLJ10052), mRNA
NM 017977	Homo sapiens hypothetical protein FLJ10044 (FLJ10044), mRNA
NM 017974	Homo sapiens hypothetical protein FLJ10040 (FLJ10040), mRNA
NM_018410	Homo sapiens hypothetical protein FLJ10035 (FLJ10035), mRNA
- 1112_010110	Homo sapiens hypothetical protein DKFZp762E1312 (DKFZp762E1312), mRNA
NM_018423	
	Homo sapiens hypothetical protein DKFZp761P1010 (DKFZp761P1010), mRNA
NM_017597	
_	Homo sapiens hypothetical protein DKFZp761K1824 (DKFZp761K1824), mRNA
NM_018422	Homo sapiens hypothetical protein DKFZp761K1423 (DKFZp761K1423),
_	mRNA
NM_018421	Homo sapiens hypothetical protein DKFZp761D1823 (DKFZp761D1823),
	mRNA
NM_017599	Homo sapiens transmembrane protein vezatin (VEZATIN), mRNA
NM_017594	Homo sapiens hypothetical protein DKFZp761C07121 (DKFZp761C07121),
	mRNA
NM_017535	Homo sapiens hypothetical protein DKFZp566H0824 (DKFZp566H0824),
	mRNA
NM_018705	Homo sapiens hypothetical protein DKFZp547G183 (DKFZp547G183), mRNA
NM_017604	Homo sapiens KIAA1023 protein (KIAA1023), mRNA
	, IIIKIVA

NM 017559	Homo sapiens hypothetical protein DKFZp434H2215 (DKFZp434H2215),
_	mRNA
NM_017598	Homo sapiens hypothetical protein DKFZp434C0923 (DKFZp434C0923), mRNA
NM_017577	Homo sapiens hypothetical protein DKFZp434C0328 (DKFZp434C0328), mRNA
NM_014612	Homo sapiens C9orf10 protein (C9orf10), mRNA
NM_018460	Homo sapiens uncharacterized bone marrow protein BM046 (BM046), mRNA
NM_018459	Homo sapiens uncharacterized bone marrow protein BM045 (BM045), mRNA
NM_018451	Homo sapiens centrosomal P4.1-associated protein (CPAP), mRNA
NM_018450	Homo sapiens uncharacterized bone marrow protein BM029 (BM029), mRNA
NM_018674	Homo sapiens putative acid-sensing ion channel (ASIC4), mRNA
NM_017435	Homo sapiens solute carrier family 21 (organic anion transporter), member 14 (SLC21A14), mRNA
NM_016848	Homo sapiens neuronal Shc (SHC3), mRNA
NM_017432	Homo sapiens prostate tumor over expressed gene 1 (PTOV1), mRNA
NM_016953	Homo sapiens phosphodiesterase 11A (PDE11A), mRNA
NM_013242	Homo sapiens similar to mouse Glt3 or D. malanogaster transcription factor IIB (AF093680), mRNA
NM 016267	Homo sapiens TONDU (TONDU), mRNA
NM_015859	Homo sapiens general transcription factor IIA, 1 (37kD and 19kD subunits) (GTF2A1), mRNA
NM 016271	Homo sapiens STRIN protein (STRIN), mRNA
NM 016584	Homo sapiens interleukin 23, alpha subunit p19 (IL23A), mRNA
NM 016329	Homo sapiens RU1 (RU1), mRNA
NM_016337	Homo sapiens RNB6 (RNB6), mRNA
NM_016146	Homo sapiens PTD009 protein (PTD009), mRNA
NM_016145	Homo sapiens PTD008 protein (PTD008), mRNA
NM_016144	Homo sapiens PTD002 protein (PTD002), mRNA
NM_016147	Homo sapiens protein phosphatase methylesterase-1 (PME-1), mRNA
NM_016445	Homo sapiens pleckstrin 2 (mouse) homolog (PLEK2), mRNA
NM_016170	Homo sapiens NCX protein (NCX), mRNA
NM_016132	Homo sapiens myelin gene expression factor 2 (MEF-2), mRNA
NM_016586	Homo sapiens MBIP protein (MBIP), mRNA
NM_016547	Homo sapiens calcium binding protein Cab45 precursor (Cab45), mRNA
NM_016530	Homo sapiens RAB-8b protein (LOC51762), mRNA
NM_016442	Homo sapiens type 1 tumor necrosis factor receptor shedding aminopeptidase regulator (ARTS-1), mRNA
NM_016438	Homo sapiens CLST 11240 protein (CLST11240), mRNA
NM_016340	Homo sapiens rap guanine nucleotide exchange factor (RA-GEF-2), mRNA
NM_016306	Homo sapiens DnaJ (Hsp40) homolog, subfamily B, member 11 (DNAJB11), mRNA
NM_016292	Homo sapiens heat shock protein 75 (TRAP1), mRNA
NM_016248	Homo sapiens A kinase (PRKA) anchor protein 11 (AKAP11), mRNA
NM_016207	Homo sapiens cleavage and polyadenylation specific factor 3, 73kD subunit (CPSF3), mRNA
NM_016163	Homo sapiens vesicle transport-related protein (RA410), mRNA
NM_016106	Homo sapiens vesicle transport-related protein (RA410), mRNA
NM_016081	Homo sapiens palladin (KIAA0992), mRNA
NM_015934	Homo sapiens nucleolar protein NOP5/NOP58 (NOP5/NOP58), mRNA
NM_015925	Homo sapiens liver-specific bHLH-Zip transcription factor (LISCH7), mRNA
NM 015878	Homo sapiens ornithine decarboxylase antizyme inhibitor (OAZIN), mRNA

NM_016284	MARIO ALLO ALLO ALLO ALLO ALLO ALLO ALLO AL
NM_016645	Homo sapiens mesenchymal stem cell protein DSC02 (NELICETED - PALA
NM_016631	110mo sapiens chromosome 21 open reading frame 66 (C21orf66) DNA
NM_016576	fromo sapiens GMPRZ for guanosine mononhosphate reductors isolar
) The O1 6701	(EOC51292), IIIRNA
NM_016501	Homo sapiens hypothetical protein FLJ10597 (FLJ10597), mRNA
NM_016500	Tiomo sapiens hypothetical protein (LOC51260) mPNA
NM_016487	Homo sapiens HSPC230 gene (HSPC230) mRNA
NM_016480	riomo sapiens PABP-interacting protein 2 (PAIP2) mPNIA
NM_016433	Homo sapiens glycolipid transfer protein (GITD) mDNIA
NM_016369	Homo sapiens claudin 18 (CLDN18) mRNA
NM_016359	Homo sapiens nucleolar protein ANKT (ANKT) mDNA
NM_016246	nomo sapiens retinal short-chain dehydrogenase/reductase retopp?
ND 6 016106	(EOC31171), IIRNA
NM_016186	Homo sapiens serine (or cysteine) proteinase inhibitor, clade A (alpha-1
ND ( 016100	unuproteniase, anguypsin), member 10 (SERPINIA 10)
NM_016180	Tionio sapiens Alivi-1 protein (MATP) mRNA
NM_016176	Homo sapiens calcium binding protein Cab45 pregursor (Cab45) DNA
NM_016174	Tionio sapiens cerebral cell adhesion molecule (I OC51140) DATA
NM_016131	Tiolio Sapiciis RABIU, member RAS oncogene femily (DADIO)
NM_016031	Tromo sapicits ciongation of very long chain fatty saids (EEXII /EI a grap term
NM 015055	1 James 1 (DEO VEI), IIIKINA
NM_015955	Homo sapiens C21orf19-like protein (LOC51072), mRNA
NM_015931	Homo sapiens fis485 (LOC51066), mRNA
NM_015879	Homo sapiens sialyltransferase 8C (alpha2,3Galbeta1,4GlcNAcalpha 2,8-
NM_016368	Total filansiciase) (SIATSC), mRNA
NM_016488	Homo sapiens myo-inositol 1-phosphate synthase A1 (ISYNA1), mRNA
NM_016478	Tromo sapiens hypothetical profein (HSPC232) mDNA
NM_016463	Homo sapiens hypothetical protein (HSPC216), mRNA
NM_016410	Homo sapiens hypothetical protein (HSPC195), mRNA
NM_016406	Homo sapiens hypothetical protein HSPC177 (HSPC177), mRNA
NM_016401	Homo sapiens hypothetical protein (HSPC155), mRNA
NM_016400	Homo sapiens hypothetical protein (HSPC138), mRNA
NM_016396	Homo sapiens Huntingtin interacting protein K (HYPK), mRNA
NM_016391	Homo sapiens hypothetical protein (HSPC129), mRNA
NM 015933	Homo sapiens hypothetical protein (HSPC111), mRNA
NM_015932	Homo sapiens hypothetical protein (HSPC016), mRNA
NM_016172	Homo sapiens hypothetical protein (HSPC014), mRNA
	Homo sapiens putative glialblastoma cell differentiation-related (GDBR1), mRNA
NM_016194	114411
	Homo sapiens guanine nucleotide binding protein (G protein), beta 5 (GNB5), mRNA
NM_016196	
NM_016553	Homo sapiens KIAA0682 gene product (KIAA0682), mRNA
NM_016195	Homo sapiens Muchae phone I ho
NM_016550	Homo sapiens M-phase phosphoprotein 1 (MPHOSPH1), mRNA
	Homo sapiens HeLa cyclin-dependent kinase 2 interacting protein (CINP), mRNA
NM_016623	111111111111111111111111111111111111111
NM_016237	Homo sapiens hypothetical protein (BM-009), mRNA
NM_016108	Homo sapiens anaphase promoting complex subunit 5 (ANAPC5), mRNA
NM_014886	Homo sapiens androgen induced protein (AIG-1), mRNA  Homo sapiens hypothetical protein (AIG-1), mRNA
NM_014035	Homo sapiens SBRI31 protein (YR-29), mRNA Homo sapiens SBRI31 protein (SBRI31)
	Homo sapiens SBBI31 protein (SBBI31), mRNA

NM_014868	Homo sapiens ring finger protein 10 (RNF10), mRNA
NM_014092	Homo sapiens PRO1575 protein (PRO1575), mRNA
NM_014138	Homo sapiens PRO0659 protein (PRO0659), mRNA
NM_014135	Homo sapiens PRO0641 protein (PRO0641), mRNA
NM_014134	Homo sapiens PRO0628 protein (PRO0628), mRNA
NM_014133	Homo sapiens PRO0618 protein (PRO0618), mRNA
NM_014076	Homo sapiens PRO0611 protein (PRO0611), mRNA
NM_014074	Homo sapiens PRO0529 protein (PRO0529), mRNA
NM_014129	Homo sapiens PRO0478 protein (PRO0478), mRNA
NM_014126	Homo sapiens PRO0365 protein (PRO0365), mRNA
NM_014124	Homo sapiens PRO0255 protein (PRO0255), mRNA
NM_014121	Homo sapiens PRO0233 protein (PRO0233), mRNA
NM_014120	Homo sapiens PRO0214 protein (PRO0214), mRNA
NM_014118	Homo sapiens PRO0159 protein (PRO0159), mRNA
NM_014117	Homo sapiens PRO0149 protein (PRO0149), mRNA
NM_014116	Homo sapiens PRO0132 protein (PRO0132), mRNA
NM_015364	Homo sapiens MD-2 protein (MD-2), mRNA
NM_014020	Homo sapiens LR8 protein (LR8), mRNA
NM_014931	Homo sapiens KIAA1115 protein (KIAA1115), mRNA
NM_014901	Homo sapiens KIAA1100 protein (KIAA1100), mRNA
NM_014908	Homo sapiens KIAA1094 protein (KIAA1094), mRNA
NM_014906	Homo sapiens KIAA1072 protein (KIAA1072), mRNA
NM_014932	Homo sapiens neuroligin 1 (NLGN1), mRNA
NM_014894	Homo sapiens KIAA1056 protein (KIAA1056), mRNA
NM_014956	Homo sapiens KIAA1052 protein (KIAA1052), mRNA
NM_014928	Homo sapiens KIAA1046 protein (KIAA1046), mRNA
NM_014909	Homo sapiens KIAA1036 protein (KIAA1036), mRNA
NM_014939	Homo sapiens KIAA1012 protein (KIAA1012), mRNA
NM_014895	Homo sapiens KIAA1009 protein (KIAA1009), mRNA
NM_014960	Homo sapiens KIAA1001 protein (KIAA1001), mRNA
NM_014950	Homo sapiens KIAA0997 protein (KIAA0997), mRNA
NM_014934	Homo sapiens zinc-finger protein DZIP1 (DZIP1), mRNA
NM_014023	Homo sapiens KIAA0982 protein (KIAA0982), mRNA
NM_014900	Homo sapiens KIAA0977 protein (KIAA0977), mRNA
NM_014929	Homo sapiens KIAA0971 protein (KIAA0971), mRNA
NM_014935	Homo sapiens phosphoinositol 3-phosphate-binding protein-2 (PEPP3), mRNA
NM_014937	Homo sapiens Sac domain-containing inositol phosphatase 2 (SAC2), mRNA
NM_014902	Homo sapiens KIAA0964 protein (KIAA0964), mRNA
NM_014898	Homo sapiens KIAA0961 protein (KIAA0961), mRNA
NM_014942	Homo sapiens ankyrin repeat domain 6 (ANKRD6), mRNA
NM_014959	Homo sapiens tumor up-regulated CARD-containing antagonist of caspase nine
	(TUCAN), mRNA
NM_014952	Homo sapiens KIAA0945 protein (KIAA0945), mRNA
NM_014904	Homo sapiens KIAA0941 protein (Rab11-FIP2), mRNA
NM_014903	Homo sapiens KIAA0938 protein (KIAA0938), mRNA
NM_014897	Homo sapiens KIAA0924 protein (KIAA0924), mRNA
NM_014883	Homo sapiens KIAA0914 gene product (KIAA0914), mRNA
NM_014949	Homo sapiens KIAA0907 protein (KIAA0907), mRNA
NM_014896	Homo sapiens KIAA0894 protein (KIAA0894), mRNA
NM_014969	Homo sapiens KIAA0893 protein (KIAA0893), mRNA
NM_014966	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 30 (DDX30),
	mRNA

37.5.05.55	
NM_015377	Homo sapiens KIAA0889 protein (KIAA0889), mRNA
NM_014936	Homo sapiens ectonucleotide pyrophosphatase/phosphodiesterase 4 (putative
	function) (ENPP4), mRNA
NM_014940	Homo sapiens KIAA0872 protein (KIAA0872), mRNA
NM_014943	Homo sapiens KIAA0854 protein (KIAA0854), mRNA
NM_014926	Homo sapiens KIAA0848 protein (KIAA0848), mRNA
NM_014945	Homo sapiens KIAA0843 protein (KIAA0843), mRNA
NM_014924	Homo sapiens KIAA0831 protein (KIAA0831), mRNA
NM_014703	Homo sapiens KIAA0800 gene product (KIAA0800), mRNA
NM_014650	Homo sapiens KIAA0798 gene product (KIAA0798), mRNA
NM_014660	Homo sapiens KIAA0783 gene product (KIAA0783), mRNA
NM_014726	Homo sapiens KIAA0775 gene product (KIAA0775), mRNA
NM_014690	Homo sapiens KIAA0773 gene product (KIAA0773), mRNA
NM_014805	Homo sapiens KIAA0766 gene product (KIAA0766), mRNA
NM_014869	Homo sapiens KIAA0763 gene product (KIAA0763), mRNA
NM_014804	Homo sapiens KIAA0753 gene product (KIAA0753), mRNA
NM_014632	Homo sapiens KIAA0750 gene product (KIAA0750), mRNA
NM_014796	Homo sapiens KIAA0748 gene product (KIAA0748), mRNA
NM_014719	Homo sapiens KIAA0738 gene product (KIAA0738), mRNA
NM_014828	Homo sapiens KIAA0737 gene product (KIAA0737), mRNA
NM_014849	Homo sapiens likely ortholog of mouse synaptic vesicle glycoprotein 2a (SV2),
NB/ 01/040	mRNA
NM_014848	Homo sapiens synaptic vesicle protein 2B homolog (SV2B), mRNA
NM_014718	Homo sapiens KIAA0726 gene product (KIAA0726), mRNA
NM_014652	Homo sapiens importin 13 (IMP13), mRNA
NM_014867	Homo sapiens KIAA0711 gene product (KIAA0711), mRNA
NM_014852	Homo sapiens KIAA0682 gene product (KIAA0682), mRNA
NM_014663	Homo sapiens KIAA0677 gene product (KIAA0677), mRNA
NM_014648 NM_014779	Homo sapiens KIAA0675 gene product (KIAA0675), mRNA
NM 014811	Homo sapiens KIAA0669 gene product (KIAA0669), mRNA
NM 014817	Homo sapiens KIAA0649 gene product (KIAA0649), mRNA
NM 015046	Homo sapiens KIAA0644 gene product (KIAA0644), mRNA
NM 014694	Homo sapiens KIAA0625 protein (KIAA0625), mRNA
NM 014832	Homo sapiens KIAA0605 gene product (KIAA0605), mRNA
NM 014749	Homo sapiens KIAA0603 gene product (KIAA0603), mRNA
NM_014668	Homo sapiens KIAA0586 gene product (KIAA0586), mRNA
NM_014709	Homo sapiens KIAA0575 gene product (KIAA0575), mRNA
NM 014704	Homo sapiens KIAA0570 gene product (KIAA0570), mRNA
NM_014790	Homo sapiens KIAA0562 gene product (KIAA0562), mRNA
NM 014731	Homo sapiens KIAA0555 gene product (KIAA0555), mRNA
NM 014793	Homo sapiens KIA 40547 gene product (KIA 40547), mRNA
NM_014825	Homo sapiens KIAA0547 gene product (KIAA0547), mRNA
NM 014840	Homo sapiens KIA A0527 come and a (KIA A0527) PNA
NM 014682	Homo sapiens KIAA0537 gene product (KIAA0537), mRNA
NM 014851	Homo sapiens KIA 00460 gene product (KIA 00460), mRNA
NM 014638	Homo sapiens KIAA0469 gene product (KIAA0469), mRNA
NM 015556	Homo sapiens KIAA0450 gene product (KIAA0450), mRNA
NM_014801	Homo sapiens KIAA0440 protein (KIAA0440), mRNA
NM 014772	Homo sapiens KIAA0435 gene product (KIAA0435), mRNA
NM_014631	Homo sapiens KIAA0427 gene product (KIAA0427), mRNA
NM 014702	Homo sapiens KIAA0418 gene product (KIAA0418), mRNA
~ 1111_UIT/UZ	Homo sapiens KIAA0408 gene product (KIAA0408), mRNA

NM 014672 Homo sapiens KIAA0391 gene product (KIAA0391), mRNA NM 014717 Homo sapiens KIAA0390 gene product (KIAA0390), mRNA NM 014872 Homo sapiens KIAA0355 gene product (KIAA0355), mRNA NM 014830 Homo sapiens KIAA0352 gene product (KIAA0354), mRNA NM 014830 Homo sapiens KIAA0352 gene product (KIAA0354), mRNA NM 014636 Homo sapiens KIAA0352 gene product (KIAA0354), mRNA NM 014636 Homo sapiens KIAA0352 gene product (KIAA0353), mRNA NM 014637 Homo sapiens KIAA0336 gene product (KIAA0336), mRNA NM 014803 Homo sapiens KIAA0336 gene product (KIAA0339), mRNA NM 014803 Homo sapiens KIAA0339 gene product (KIAA0339), mRNA NM 014821 Homo sapiens KIAA0317 gene product (KIAA0317), mRNA NM 014821 Homo sapiens KIAA0317 gene product (KIAA0317), mRNA NM 014791 Homo sapiens KIAA02596 gene product (KIAA0296), mRNA NM 014734 Homo sapiens KIAA02269 gene product (KIAA0247), mRNA NM 014760 Homo sapiens KIAA0218 gene product (KIAA0218), mRNA NM 014760 Homo sapiens KIAA0218 gene product (KIAA0218), mRNA NM 014735 Homo sapiens KIAA0211 gene product (KIAA0211), mRNA NM 014744 Homo sapiens KIAA0211 gene product (KIAA0211), mRNA NM 014744 Homo sapiens KIAA0219 gene product (KIAA0211), mRNA NM 014744 Homo sapiens KIAA0219 gene product (KIAA0211), mRNA NM 014744 Homo sapiens KIAA0219 gene product (KIAA0210), mRNA NM 014744 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA NM 014751 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA NM 014760 Homo sapiens KIAA0189 gene product (KIAA0189), mRNA NM 014761 Homo sapiens KIAA0187 gene product (KIAA0189), mRNA NM 014791 Homo sapiens KIAA0187 gene product (KIAA0189), mRNA NM 014794 Homo sapiens KIAA0187 gene product (KIAA0189), mRNA NM 014794 Homo sapiens KIAA0189 gene product (KIAA0189), mRNA NM 014760 Homo sapiens KIAA0119 gene product (KIAA0118), mRNA NM 014760 Homo sapiens KIAA0119 gene product (KIAA0118), mRNA NM 014760 Homo sapiens KIAA0119 gene product (KIAA0189), mRNA NM 014794 Homo sapiens KIAA0119 gene product (KIAA0019), mRNA NM 014794 Homo sapiens KIAA0119 gene product (KIAA0019), mRN
NM 014820 Homo sapiens KIAA0355 gene product (KIAA0355), mRNA NM 014830 Homo sapiens KIAA0354 gene product (KIAA0352), mRNA NM 014830 Homo sapiens RIAA0352 gene product (KIAA0352), mRNA NM 014830 Homo sapiens Ral guanine nucleotide exchange factor RalGPS1A (RALGPS1A), mRNA NM 014831 Homo sapiens RIAA0335 gene product (KIAA0336), mRNA NM 014803 Homo sapiens KIAA0335 gene product (KIAA0336), mRNA NM 014803 Homo sapiens KIAA0335 gene product (KIAA0335), mRNA NM 014844 Homo sapiens KIAA0329 gene product (KIAA0329), mRNA NM 014821 Homo sapiens KIAA0317 gene product (KIAA0329), mRNA NM 014899 Homo sapiens KIAA0296 gene product (KIAA0296), mRNA NM 014742 Homo sapiens KIAA0246 gene product (KIAA0255), mRNA NM 014743 Homo sapiens KIAA0247 gene product (KIAA0247), mRNA NM 014760 Homo sapiens KIAA0218 gene product (KIAA0247), mRNA NM 014760 Homo sapiens KIAA0218 gene product (KIAA0218), mRNA NM 014760 Homo sapiens KIAA0218 gene product (KIAA0218), mRNA NM 014760 Homo sapiens KIAA0218 gene product (KIAA0218), mRNA NM 014760 Homo sapiens KIAA0218 gene product (KIAA0218), mRNA NM 014760 Homo sapiens KIAA0218 gene product (KIAA0211), mRNA NM 014760 Homo sapiens KIAA0218 gene product (KIAA0211), mRNA NM 014760 Homo sapiens KIAA0187 gene product (KIAA0210), mRNA NM 014760 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA NM 014760 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA NM 014760 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA NM 014760 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA NM 014760 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA NM 014760 Homo sapiens KIAA0188 gene product (KIAA0187), mRNA NM 014760 Homo sapiens KIAA0187 gene product (KIAA0161), mRNA NM 014630 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA NM 014773 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA NM 014773 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA NM 014774 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA NM 014774 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA NM 014760 Homo sapiens KIAA0188 gene
NM 014872 Homo sapiens KIAA0354 gene product (KIAA0354), mRNA NM 014830 Homo sapiens RIAA0352 gene product (KIAA0352), mRNA NM 014636 Homo sapiens RIAA0353 gene product (KIAA0336), mRNA NM 014635 Homo sapiens KIAA0336 gene product (KIAA0336), mRNA NM 014803 Homo sapiens KIAA0335 gene product (KIAA0335), mRNA NM 014844 Homo sapiens KIAA0339 gene product (KIAA0335), mRNA NM 014841 Homo sapiens KIAA0317 gene product (KIAA0317), mRNA NM 014821 Homo sapiens KIAA0329 gene product (KIAA0317), mRNA NM 014694 Homo sapiens KIAA0296 gene product (KIAA0296), mRNA NM 014742 Homo sapiens KIAA0247 gene product (KIAA0295), mRNA NM 014743 Homo sapiens KIAA0247 gene product (KIAA0247), mRNA NM 014745 Homo sapiens KIAA0218 gene product (KIAA0218), mRNA NM 014735 Homo sapiens KIAA0211 gene product (KIAA0218), mRNA NM 014744 Homo sapiens KIAA0211 gene product (KIAA0211), mRNA NM 014745 Homo sapiens KIAA0210 gene product (KIAA0210), mRNA NM 014745 Homo sapiens KIAA0210 gene product (KIAA0189), mRNA NM 014753 Homo sapiens KIAA0189 gene product (KIAA0189), mRNA NM 014753 Homo sapiens KIAA0189 gene product (KIAA0189), mRNA NM 014753 Homo sapiens KIAA0189 gene product (KIAA0189), mRNA NM 014764 Homo sapiens KIAA0189 gene product (KIAA0189), mRNA NM 014753 Homo sapiens KIAA0189 gene product (KIAA0189), mRNA NM 014754 Homo sapiens KIAA0118 gene product (KIAA0189), mRNA NM 014754 Homo sapiens KIAA0116 gene product (KIAA0189), mRNA NM 014764 Homo sapiens KIAA0116 gene product (KIAA0161), mRNA NM 01476 Homo sapiens KIAA0118 gene product (KIAA0151), mRNA NM 01479 Homo sapiens KIAA0118 gene product (KIAA0141), mRNA NM 01479 Homo sapiens KIAA0119 gene product (KIAA0111), mRNA NM 01479 Homo sapiens KIAA0119 gene product (KIAA0189), mRNA NM 01479 Homo sapiens KIAA0119 gene product (KIAA0189), mRNA NM 01479 Homo sapiens KIAA0193 gene product (KIAA0189), mRNA NM 01479 Homo sapiens KIAA0193 gene product (KIAA0195), mRNA NM 01479 Homo sapiens KIAA0195 gene product (KIAA0195), mRNA NM 01476 Homo sapiens KIAA0195 gene product (KIAA0095), mRNA NM 01476
NM 014830   Homo sapiens KIAA0352 gene product (KIAA0352), mRNA
NM_014636 Homo sapiens Ral guanine nucleotide exchange factor RalGPS1A (RALGPS1A), mRNA  NM_014803 Homo sapiens KIAA0336 gene product (KIAA0336), mRNA  NM_014803 Homo sapiens KIAA0335 gene product (KIAA0335), mRNA  NM_014844 Homo sapiens KIAA0329 gene product (KIAA0317), mRNA  NM_014821 Homo sapiens KIAA0317 gene product (KIAA0317), mRNA  NM_014699 Homo sapiens KIAA0296 gene product (KIAA0296), mRNA  NM_014742 Homo sapiens KIAA0255 gene product (KIAA0296), mRNA  NM_014743 Homo sapiens KIAA0247 gene product (KIAA0247), mRNA  NM_014734 Homo sapiens KIAA0218 gene product (KIAA0218), mRNA  NM_014735 Homo sapiens KIAA0218 gene product (KIAA0211), mRNA  NM_014735 Homo sapiens KIAA0210 gene product (KIAA0211), mRNA  NM_014745 Homo sapiens KIAA0210 gene product (KIAA0210), mRNA  NM_014751 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA  NM_014753 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA  NM_014754 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA  NM_014751 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA  NM_014764 Homo sapiens KIAA0161 gene product (KIAA0161), mRNA  NM_014764 Homo sapiens KIAA0161 gene product (KIAA0161), mRNA  NM_014764 Homo sapiens KIAA0187 gene product (KIAA0161), mRNA  NM_014764 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA  NM_014773 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA  NM_014773 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA  NM_014774 Homo sapiens KIAA0187 gene product (KIAA0141), mRNA  NM_014794 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA  NM_014796 Homo sapiens KIAA0187 gene product (KIAA00187), mRNA  NM_
CRALGPS1A), mRNA
(RALGPS1A), mRNA  NM 014803 Homo sapiens KIAA0336 gene product (KIAA0336), mRNA  NM 014844 Homo sapiens KIAA0335 gene product (KIAA0329), mRNA  NM 014844 Homo sapiens KIAA0317 gene product (KIAA0329), mRNA  NM 014821 Homo sapiens KIAA0296 gene product (KIAA0317), mRNA  NM 014821 Homo sapiens KIAA0296 gene product (KIAA0296), mRNA  NM 014742 Homo sapiens KIAA0296 gene product (KIAA0296), mRNA  NM 014734 Homo sapiens KIAA0257 gene product (KIAA0247), mRNA  NM 014734 Homo sapiens KIAA0218 gene product (KIAA0247), mRNA  NM 014736 Homo sapiens KIAA0218 gene product (KIAA0218), mRNA  NM 014630 Homo sapiens KIAA0216 gene product (KIAA0211), mRNA  NM 014734 Homo sapiens KIAA0210 gene product (KIAA0211), mRNA  NM 014734 Homo sapiens KIAA0210 gene product (KIAA0210), mRNA  NM 014735 Homo sapiens KIAA0187 gene product (KIAA0189), mRNA  NM 014791 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA  NM 014791 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA  NM 014791 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA  NM 014791 Homo sapiens KIAA0161 gene product (KIAA0161), mRNA  NM 014791 Homo sapiens KIAA0161 gene product (KIAA0161), mRNA  NM 014794 Homo sapiens KIAA0161 gene product (KIAA0161), mRNA  NM 014902 Homo sapiens KIAA0161 gene product (KIAA0161), mRNA  NM 014940 Homo sapiens KIAA0118 gene product (KIAA0144), mRNA  NM 014940 Homo sapiens KIAA0118 gene product (KIAA0144), mRNA  NM 014791 Homo sapiens KIAA0118 gene product (KIAA0118), mRNA  NM 014792 Homo sapiens KIAA0118 gene product (KIAA0118), mRNA  NM 014794 Homo sapiens KIAA0118 gene product (KIAA0118), mRNA  NM 014794 Homo sapiens KIAA0118 gene product (KIAA0118), mRNA  NM 014794 Homo sapiens KIAA0138 gene product (KIAA0118), mRNA  NM 014794 Homo sapiens KIAA0138 gene product (KIAA0118), mRNA  NM 014794 Homo sapiens KIAA0138 gene product (KIAA0193), mRNA  NM 014796 Homo sapiens KIAA0187 gene product (KIAA0193), mRNA  NM 014796 Homo sapiens KIAA0187 gene product (KIAA0193), mRNA  NM 014796 Homo sapiens KIAA0187 gene product (KIAA0193), mRNA  NM 014769 Ho
NM 014635 Homo sapiens KIAA0336 gene product (KIAA0336), mRNA NM 014844 Homo sapiens KIAA0335 gene product (KIAA0335), mRNA NM 014844 Homo sapiens KIAA0329 gene product (KIAA0329), mRNA NM 014821 Homo sapiens KIAA0317 gene product (KIAA0329), mRNA NM 014891 Homo sapiens KIAA0296 gene product (KIAA0296), mRNA NM 014742 Homo sapiens KIAA0295 gene product (KIAA0296), mRNA NM 014734 Homo sapiens KIAA0255 gene product (KIAA0247), mRNA NM 014760 Homo sapiens KIAA0218 gene product (KIAA0247), mRNA NM 014735 Homo sapiens KIAA0218 gene product (KIAA0218), mRNA NM 014740 Homo sapiens KIAA0211 gene product (KIAA0211), mRNA NM 014745 Homo sapiens KIAA0210 gene product (KIAA0211), mRNA NM 014745 Homo sapiens KIAA0210 gene product (KIAA0210), mRNA NM 014745 Homo sapiens KIAA0189 gene product (KIAA0210), mRNA NM 014753 Homo sapiens KIAA0187 gene product (KIAA0210), mRNA NM 014761 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA NM 014764 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA NM 014764 Homo sapiens KIAA0161 gene product (KIAA0161), mRNA NM 014847 Homo sapiens KIAA0161 gene product (KIAA0161), mRNA NM 01402 Homo sapiens KIAA0144 gene product (KIAA0144), mRNA NM 014847 Homo sapiens KIAA0144 gene product (KIAA0144), mRNA NM 014792 Homo sapiens KIAA0141 gene product (KIAA0141), mRNA NM 014792 Homo sapiens KIAA0118 gene product (KIAA0141), mRNA NM 014794 Homo sapiens KIAA0118 gene product (KIAA0141), mRNA NM 014792 Homo sapiens KIAA0118 gene product (KIAA0138), mRNA NM 014794 Homo sapiens KIAA0119 gene product (KIAA0118), mRNA NM 014796 Homo sapiens KIAA0119 gene product (KIAA0141), mRNA NM 014797 Homo sapiens KIAA0199 gene product (KIAA0118), mRNA NM 014799 Homo sapiens KIAA0199 gene product (KIAA0103), mRNA NM 014760 Homo sapiens KIAA0199 gene product (KIAA0110), mRNA NM 014761 Homo sapiens KIAA0099 gene product (KIAA0099), mRNA NM 014761 Homo sapiens KIAA0099 gene product (KIAA0099), mRNA NM 014761 Homo sapiens KIAA0099 gene product (KIAA0099), mRNA NM 014871 Homo sapiens KIAA0099 gene product (KIAA0099), mRNA N
NM_014803 Homo sapiens KIAA0335 gene product (KIAA0335), mRNA NM_014844 Homo sapiens KIAA0329 gene product (KIAA0329), mRNA NM_014821 Homo sapiens KIAA0317 gene product (KIAA0317), mRNA NM_014699 Homo sapiens KIAA0296 gene product (KIAA0296), mRNA NM_014742 Homo sapiens KIAA0255 gene product (KIAA0255), mRNA NM_014744 Homo sapiens KIAA0247 gene product (KIAA0255), mRNA NM_014734 Homo sapiens KIAA0218 gene product (KIAA0218), mRNA NM_014735 Homo sapiens KIAA0218 gene product (KIAA0218), mRNA NM_014735 Homo sapiens KIAA0218 gene product (KIAA0211), mRNA NM_014630 Homo sapiens KIAA0211 gene product (KIAA0210), mRNA NM_014744 Homo sapiens KIAA0210 gene product (KIAA0210), mRNA NM_014745 Homo sapiens KIAA0189 gene product (KIAA0189), mRNA NM_014751 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA NM_014791 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA NM_014633 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA NM_014634 Homo sapiens KIAA0161 gene product (KIAA0161), mRNA NM_014634 Homo sapiens KIAA0155 gene product (KIAA0155), mRNA NM_014604 Homo sapiens KIAA0141 gene product (KIAA0144), mRNA NM_014649 Homo sapiens KIAA0144 gene product (KIAA0144), mRNA NM_014649 Homo sapiens KIAA0143 gene product (KIAA0144), mRNA NM_01469 Homo sapiens KIAA0118 protein (KIAA0118), mRNA NM_01469 Homo sapiens KIAA0118 protein (KIAA0118), mRNA NM_014791 Homo sapiens KIAA0118 gene product (KIAA0111), mRNA NM_014691 Homo sapiens KIAA0118 gene product (KIAA0111), mRNA NM_014699 Homo sapiens KIAA0119 gene product (KIAA0111), mRNA NM_014690 Homo sapiens KIAA0119 gene product (KIAA0119), mRNA NM_014691 Homo sapiens KIAA0019 gene product (KIAA0119), mRNA NM_014693 Homo sapiens KIAA0019 gene product (KIAA0119), mRNA NM_014694 Homo sapiens KIAA0019 gene product (KIAA0199), mRNA NM_014699 Homo sapiens KIAA0092 gene product (KIAA0099), mRNA NM_014690 Homo sapiens KIAA0092 gene product (KIAA0099), mRNA NM_014691 Homo sapiens KIAA0092 gene product (KIAA0099), mRNA NM_014691 Homo sapiens KIAA0092 gene product (KIAA0099), mRNA NM_014691 Ho
NM 014844 Homo sapiens KIAA0329 gene product (KIAA0329), mRNA NM 014821 Homo sapiens KIAA0317 gene product (KIAA0317), mRNA NM 014699 Homo sapiens KIAA0296 gene product (KIAA0296), mRNA NM 014742 Homo sapiens KIAA0295 gene product (KIAA0295), mRNA NM 014734 Homo sapiens KIAA0247 gene product (KIAA0247), mRNA NM 014736 Homo sapiens KIAA0218 gene product (KIAA0218), mRNA NM 014735 Homo sapiens KIAA0218 gene product (KIAA0218), mRNA NM 014630 Homo sapiens KIAA0211 gene product (KIAA0211), mRNA NM 014744 Homo sapiens KIAA0210 gene product (KIAA0211), mRNA NM 014753 Homo sapiens KIAA0189 gene product (KIAA0189), mRNA NM 014791 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA NM 014791 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA NM 014746 Homo sapiens KIAA0161 gene product (KIAA0161), mRNA NM 01433 Homo sapiens KIAA0155 gene product (KIAA0161), mRNA NM 014002 Homo sapiens KIAA0155 gene product (KIAA0161), mRNA NM 014002 Homo sapiens KIAA0149 gene product (KIAA0144), mRNA NM 014773 Homo sapiens KIAA0149 gene product (KIAA0144), mRNA NM 014773 Homo sapiens KIAA0149 gene product (KIAA0141), mRNA NM 014774 Homo sapiens KIAA0149 gene product (KIAA0141), mRNA NM 014773 Homo sapiens KIAA0149 gene product (KIAA0141), mRNA NM 014792 Homo sapiens KIAA0149 gene product (KIAA0141), mRNA NM 014794 Homo sapiens KIAA0149 gene product (KIAA0141), mRNA NM 014794 Homo sapiens KIAA0189 gene product (KIAA0138), mRNA NM 014794 Homo sapiens KIAA0189 gene product (KIAA0138), mRNA NM 014794 Homo sapiens KIAA0189 gene product (KIAA0118), mRNA NM 014794 Homo sapiens KIAA0125 gene product (KIAA0118), mRNA NM 014796 Homo sapiens KIAA0190 gene product (KIAA0111), mRNA NM 014796 Homo sapiens KIAA0090 gene product (KIAA0090), mRNA NM 014769 Homo sapiens KIAA0090 gene product (KIAA0090), mRNA NM 014679 Homo sapiens KIAA0090 gene product (KIAA0090), mRNA NM 014679 Homo sapiens KIAA0090 gene product (KIAA0090), mRNA NM 014761 Homo sapiens KIAA0090 gene product (KIAA0090), mRNA NM 014871 Homo sapiens KIAA0090 gene product (KIAA0090), mRNA N
NM 014821 Homo sapiens KIAA0317 gene product (KIAA0317), mRNA NM 014699 Homo sapiens KIAA0296 gene product (KIAA0296), mRNA NM 014742 Homo sapiens KIAA0255 gene product (KIAA0255), mRNA NM 014734 Homo sapiens KIAA0247 gene product (KIAA0247), mRNA NM 014760 Homo sapiens KIAA0218 gene product (KIAA0218), mRNA NM 014760 Homo sapiens KIAA0218 gene product (KIAA0215), mRNA NM 014735 Homo sapiens KIAA0215 gene product (KIAA0211), mRNA NM 014744 Homo sapiens KIAA0210 gene product (KIAA0211), mRNA NM 014745 Homo sapiens KIAA0210 gene product (KIAA0210), mRNA NM 014753 Homo sapiens KIAA0189 gene product (KIAA0189), mRNA NM 014751 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA NM 014761 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA NM 014764 Homo sapiens KIAA0161 gene product (KIAA0161), mRNA NM 014633 Homo sapiens KIAA0165 gene product (KIAA0155), mRNA NM 014649 Homo sapiens KIAA0144 gene product (KIAA0155), mRNA NM 014847 Homo sapiens KIAA0144 gene product (KIAA0144), mRNA NM 014773 Homo sapiens KIAA0149 gene product (KIAA0144), mRNA NM 014774 Homo sapiens KIAA0138 gene product (KIAA0141), mRNA NM 014772 Homo sapiens KIAA0138 gene product (KIAA0138), mRNA NM 014773 Homo sapiens KIAA0138 gene product (KIAA0118), mRNA NM 014794 Homo sapiens KIAA0138 gene product (KIAA0111), mRNA NM 014790 Homo sapiens KIAA0113 gene product (KIAA0118), mRNA NM 014790 Homo sapiens KIAA0119 gene product (KIAA0118), mRNA NM 014790 Homo sapiens KIAA0119 gene product (KIAA0111), mRNA NM 014791 Homo sapiens KIAA0190 gene product (KIAA0103), mRNA NM 014794 Homo sapiens KIAA0190 gene product (KIAA0103), mRNA NM 014796 Homo sapiens KIAA0190 gene product (KIAA0103), mRNA NM 014791 Homo sapiens KIAA0092 gene product (KIAA0092), mRNA NM 014791 Homo sapiens KIAA0092 gene product (KIAA0095), mRNA NM 014791 Homo sapiens KIAA0092 gene product (KIAA0095), mRNA NM 014791 Homo sapiens KIAA0092 gene product (KIAA0087), mRNA NM 014791 Homo sapiens KIAA0093 gene product (KIAA0095), mRNA NM 014791 Homo sapiens KIAA0094 gene product (KIAA0095), mRNA
NM_014699 Homo sapiens KIAA0296 gene product (KIAA0296), mRNA NM_014742 Homo sapiens KIAA0255 gene product (KIAA0255), mRNA NM_014734 Homo sapiens KIAA0247 gene product (KIAA0247), mRNA NM_014735 Homo sapiens KIAA0218 gene product (KIAA0218), mRNA NM_014735 Homo sapiens KIAA0218 gene product (KIAA0218), mRNA NM_014630 Homo sapiens KIAA0211 gene product (KIAA0211), mRNA NM_014744 Homo sapiens KIAA0210 gene product (KIAA0210), mRNA NM_014753 Homo sapiens KIAA0189 gene product (KIAA0189), mRNA NM_014754 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA NM_014791 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA NM_014791 Homo sapiens KIAA0161 gene product (KIAA0161), mRNA NM_014633 Homo sapiens KIAA0161 gene product (KIAA0155), mRNA NM_014002 Homo sapiens KIAA0155 gene product (KIAA0155), mRNA NM_014002 Homo sapiens KIAA0141 gene product (KIAA0144), mRNA NM_01494 Homo sapiens KIAA0141 gene product (KIAA0144), mRNA NM_01494 Homo sapiens KIAA0141 gene product (KIAA0144), mRNA NM_014792 Homo sapiens KIAA0138 gene product (KIAA0138), mRNA NM_014794 Homo sapiens KIAA0138 gene product (KIAA0118), mRNA NM_014794 Homo sapiens KIAA0138 gene product (KIAA0118), mRNA NM_014794 Homo sapiens KIAA0118 protein (KIAA0118), mRNA NM_014790 Homo sapiens KIAA0119 gene product (KIAA0118), mRNA NM_014790 Homo sapiens KIAA0119 gene product (KIAA0110), mRNA NM_014791 Homo sapiens KIAA0103 gene product (KIAA0101), mRNA NM_014790 Homo sapiens KIAA0103 gene product (KIAA0110), mRNA NM_014791 Homo sapiens KIAA0103 gene product (KIAA0101), mRNA NM_014790 Homo sapiens KIAA0109 gene product (KIAA0101), mRNA NM_014791 Homo sapiens KIAA0091 gene product (KIAA0091), mRNA NM_014709 Homo sapiens KIAA0092 gene product (KIAA0092), mRNA NM_014709 Homo sapiens KIAA0093 gene product (KIAA0093), mRNA NM_014709 Homo sapiens KIAA0094 gene product (KIAA0094), mRNA NM_0147
NM 014742 Homo sapiens KIAA0255 gene product (KIAA0255), mRNA NM 014734 Homo sapiens KIAA0247 gene product (KIAA0247), mRNA NM 014735 Homo sapiens KIAA0218 gene product (KIAA0218), mRNA NM 014735 Homo sapiens KIAA0211 gene product (KIAA0211), mRNA NM 014630 Homo sapiens KIAA0211 gene product (KIAA0211), mRNA NM 014744 Homo sapiens KIAA0210 gene product (KIAA0210), mRNA NM 014725 Homo sapiens KIAA0189 gene product (KIAA0189), mRNA NM 014725 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA NM 014791 Homo sapiens kIAA0187 gene product (KIAA0187), mRNA NM 014791 Homo sapiens kIAA0187 gene product (KIAA0187), mRNA NM 014746 Homo sapiens KIAA0161 gene product (KIAA0161), mRNA NM 014633 Homo sapiens KIAA0155 gene product (KIAA0155), mRNA NM 014746 Homo sapiens KIAA0155 gene product (KIAA0155), mRNA NM 014701 Homo sapiens KIAA0114 gene product (KIAA0144), mRNA NM 014847 Homo sapiens KIAA0141 gene product (KIAA0141), mRNA NM 014773 Homo sapiens KIAA0118 gene product (KIAA0141), mRNA NM 014792 Homo sapiens KIAA0118 gene product (KIAA0141), mRNA NM 014792 Homo sapiens KIAA0118 gene product (KIAA0118), mRNA NM 014794 Homo sapiens KIAA0118 protein (KIAA0118), mRNA NM 014790 Homo sapiens KIAA0118 protein (KIAA0118), mRNA NM 014791 Homo sapiens KIAA0119 gene product (KIAA0111), mRNA NM 014790 Homo sapiens KIAA0103 gene product (KIAA0103), mRNA NM 014790 Homo sapiens KIAA0103 gene product (KIAA0103), mRNA NM 014791 Homo sapiens KIAA0103 gene product (KIAA0103), mRNA NM 014794 Homo sapiens KIAA0095 gene product (KIAA0095), mRNA NM 014796 Homo sapiens KIAA0095 gene product (KIAA0095), mRNA NM 014797 Homo sapiens KIAA0095 gene product (KIAA0095), mRNA NM 014799 Homo sapiens KIAA0095 gene product (KIAA0095), mRNA NM 014791 Homo sapiens KIAA0095 gene product (KIAA0095), mRNA NM 014791 Homo sapiens KIAA0095 gene product (KIAA0095), mRNA NM 014791 Homo sapiens KIAA0095 gene product (KIAA0095), mRNA NM 014791 Homo sapiens KIAA0095 gene product (KIAA0095), mRNA NM 014791 Homo sapiens KIAA0095 gene product (KIAA0095), mRNA NM 014791
NM 014734 Homo sapiens KIAA0247 gene product (KIAA0247), mRNA NM 014760 Homo sapiens KIAA0218 gene product (KIAA0218), mRNA NM 014735 Homo sapiens KIAA0215 gene product (KIAA0215), mRNA NM 014736 Homo sapiens KIAA0211 gene product (KIAA0211), mRNA NM 014744 Homo sapiens KIAA0210 gene product (KIAA0210), mRNA NM 014725 Homo sapiens KIAA02189 gene product (KIAA0189), mRNA NM 014753 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA NM 014791 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA NM 014794 Homo sapiens KIAA0161 gene product (KIAA0161), mRNA NM 014633 Homo sapiens KIAA0165 gene product (KIAA0165), mRNA NM 014634 Homo sapiens KIAA0155 gene product (KIAA0155), mRNA NM 014002 Homo sapiens KIAA0144 gene product (KIAA0144), mRNA NM 014773 Homo sapiens KIAA0141 gene product (KIAA0141), mRNA NM 014774 Homo sapiens KIAA0138 gene product (KIAA0141), mRNA NM 014792 Homo sapiens KIAA0118 gene product (KIAA0138), mRNA NM 014792 Homo sapiens KIAA0118 protein (KIAA0118), mRNA NM 014790 Homo sapiens KIAA0111 gene product (KIAA0111), mRNA NM 014790 Homo sapiens KIAA0103 gene product (KIAA0111), mRNA NM 014736 Homo sapiens KIAA0103 gene product (KIAA0101), mRNA NM 014736 Homo sapiens KIAA0109 gene product (KIAA0095), mRNA NM 014769 Homo sapiens KIAA0097 gene product (KIAA0087), mRNA NM 014760 Homo sapiens KIAA0087 gene product (KIAA0087), mRNA NM 014761 Homo sapiens KIAA0087 gene product (KIAA0087), mRNA NM 014761 Homo sapiens KIAA0087 gene product (KIAA0087), mRNA NM 014716 Homo sapiens KIAA0087 gene product (KIAA0087), mRNA NM 014716 Homo sapiens KIAA0087 gene product (KIAA0087), mRNA NM 014716 Homo sapiens KIAA0087 gene product (KIAA0087), mRNA NM 014716 Homo sapiens KIAA0087 gene product (KIAA0087), mRNA NM 014716 Homo sapiens KIAA0087 gene product (KIAA0087), mRNA NM 014716 Homo sapiens KIAA0087 gene product (KIAA0087), mRNA NM 014716 Homo sapiens KIAA0087 gene product (KIAA0087), mRNA NM 014716 Homo sapiens KIAA0087 gene product (KIAA0087), mRNA
NM 014760 Homo sapiens KIAA0218 gene product (KIAA0218), mRNA NM 014735 Homo sapiens KIAA0215 gene product (KIAA0215), mRNA NM 014630 Homo sapiens KIAA0211 gene product (KIAA0211), mRNA NM 014744 Homo sapiens KIAA0210 gene product (KIAA0210), mRNA NM 014725 Homo sapiens KIAA0189 gene product (KIAA0189), mRNA NM 014735 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA NM 014736 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA NM 014746 Homo sapiens KIAA0187 gene product (KIAA0161), mRNA NM 014631 Homo sapiens KIAA0161 gene product (KIAA0161), mRNA NM 014002 Homo sapiens KIAA0155 gene product (KIAA0155), mRNA NM 014847 Homo sapiens KIAA0144 gene product (KIAA0144), mRNA NM 014649 Homo sapiens KIAA0141 gene product (KIAA0141), mRNA NM 014649 Homo sapiens KIAA0138 gene product (KIAA0125), mRNA NM 014792 Homo sapiens KIAA0118 protein (KIAA0118), mRNA NM 014794 Homo sapiens KIAA0118 protein (KIAA0118), mRNA NM 014794 Homo sapiens KIAA0119 gene product (KIAA0111), mRNA NM 014796 Homo sapiens KIAA0101 gene product (KIAA0101), mRNA NM 014679 Homo sapiens KIAA0101 gene product (KIAA0101), mRNA NM 014679 Homo sapiens KIAA00092 gene product (KIAA0095), mRNA NM 014679 Homo sapiens KIAA0092 gene product (KIAA0095), mRNA NM 014679 Homo sapiens KIAA0092 gene product (KIAA0097), mRNA NM 014769 Homo sapiens KIAA0097 gene product (KIAA0097), mRNA NM 014769 Homo sapiens KIAA0097 gene product (KIAA0097), mRNA NM 014769 Homo sapiens KIAA0097 gene product (KIAA0097), mRNA NM 014716 Homo sapiens KIAA0087 gene product (KIAA0097), mRNA NM 014716 Homo sapiens KIAA0087 gene product (KIAA0097), mRNA NM 014716 Homo sapiens R3H domain (binds single-stranded nucleic acids) containing (R3HDM), mRNA
NM 014735 Homo sapiens KIAA0215 gene product (KIAA0215), mRNA NM 014630 Homo sapiens KIAA0211 gene product (KIAA0211), mRNA NM 014744 Homo sapiens KIAA0210 gene product (KIAA0210), mRNA NM 014725 Homo sapiens KIAA0189 gene product (KIAA0189), mRNA NM 014753 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA NM 014791 Homo sapiens likely ortholog of maternal embryonic leucine zipper kinase (KIAA0175), mRNA NM 014746 Homo sapiens KIAA0161 gene product (KIAA0161), mRNA NM 014633 Homo sapiens KIAA0155 gene product (KIAA0155), mRNA NM 014002 Homo sapiens KIAA0155 gene product (KIAA0155), mRNA NM 014847 Homo sapiens KIAA0144 gene product (KIAA0144), mRNA NM 014773 Homo sapiens KIAA0141 gene product (KIAA0141), mRNA NM 014649 Homo sapiens KIAA0138 gene product (KIAA0138), mRNA NM 014792 Homo sapiens KIAA0125 gene product (KIAA0125), mRNA NM 014794 Homo sapiens KIAA0118 protein (KIAA0118), mRNA NM 014794 Homo sapiens KIAA0110 gene product (KIAA0111), mRNA NM 014790 Homo sapiens KIAA0110 gene product (KIAA0111), mRNA NM 014791 Homo sapiens KIAA0103 gene product (KIAA0101), mRNA NM 014790 Homo sapiens KIAA0103 gene product (KIAA0101), mRNA NM 014791 Homo sapiens KIAA0103 gene product (KIAA0101), mRNA NM 014796 Homo sapiens KIAA0095 gene product (KIAA0095), mRNA NM 014669 Homo sapiens KIAA0092 gene product (KIAA0095), mRNA NM 014679 Homo sapiens KIAA0092 gene product (KIAA0097), mRNA NM 014760 Homo sapiens KIAA0092 gene product (KIAA0097), mRNA NM 014761 Homo sapiens KIAA0087 gene product (KIAA0087), mRNA NM 014716 Homo sapiens R3H domain (binds single-stranded nucleic acids) containing (R3HDM), mRNA
NM 01473 Homo sapiens KIAA0189 gene product (KIAA0187), mRNA NM 014744 Homo sapiens KIAA0189 gene product (KIAA0189), mRNA NM 014725 Homo sapiens KIAA0189 gene product (KIAA0189), mRNA NM 014753 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA NM 014791 Homo sapiens kIAA0187 gene product (KIAA0187), mRNA NM 014746 Homo sapiens KIAA0161 gene product (KIAA0161), mRNA NM 014633 Homo sapiens KIAA0165 gene product (KIAA0165), mRNA NM 014002 Homo sapiens KIAA0155 gene product (KIAA0155), mRNA NM 014047 Homo sapiens KIAA0144 gene product (KIAA0144), mRNA NM 01473 Homo sapiens KIAA0141 gene product (KIAA0141), mRNA NM 014792 Homo sapiens KIAA0138 gene product (KIAA0138), mRNA NM 014792 Homo sapiens KIAA0118 protein (KIAA0138), mRNA NM 014794 Homo sapiens KIAA0118 protein (KIAA0118), mRNA NM 01470 Homo sapiens KIAA0111 gene product (KIAA0111), mRNA NM 01473 Homo sapiens KIAA0103 gene product (KIAA0111), mRNA NM 014740 Homo sapiens KIAA0110 gene product (KIAA0103), mRNA NM 014736 Homo sapiens KIAA0103 gene product (KIAA0103), mRNA NM 014736 Homo sapiens KIAA0103 gene product (KIAA0103), mRNA NM 014669 Homo sapiens KIAA0095 gene product (KIAA0095), mRNA NM 014679 Homo sapiens KIAA0097 gene product (KIAA0097), mRNA NM 014877 Homo sapiens KIAA0087 gene product (KIAA0087), mRNA NM 014716 Homo sapiens KIAA0087 gene product (KIAA0087), mRNA NM 014716 Homo sapiens KIAA0087 gene product (KIAA0087), mRNA NM 014716 Homo sapiens KIAA0087 gene product (KIAA0087), mRNA NM 014716 Homo sapiens R3H domain (binds single-stranded nucleic acids) containing (R3HDM), mRNA
NM 014744 Homo sapiens KIAA0210 gene product (KIAA0210), mRNA NM 014725 Homo sapiens KIAA0189 gene product (KIAA0189), mRNA NM 014753 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA NM 014791 Homo sapiens kIAA0187 gene product (KIAA0187), mRNA NM 014791 Homo sapiens kIAA0161 gene product (KIAA0161), mRNA NM 014633 Homo sapiens KIAA0161 gene product (KIAA0165), mRNA NM 014002 Homo sapiens KIAA0155 gene product (KIAA0155), mRNA NM 014847 Homo sapiens KIAA0144 gene product (KIAA0144), mRNA NM 014773 Homo sapiens KIAA0144 gene product (KIAA0141), mRNA NM 014649 Homo sapiens KIAA0138 gene product (KIAA0138), mRNA NM 014792 Homo sapiens KIAA0138 gene product (KIAA0125), mRNA NM 014794 Homo sapiens KIAA0118 protein (KIAA0118), mRNA NM 014704 Homo sapiens KIAA0111 gene product (KIAA0111), mRNA NM 014704 Homo sapiens KIAA0111 gene product (KIAA0111), mRNA NM 014704 Homo sapiens KIAA0103 gene product (KIAA0101), mRNA NM 014673 Homo sapiens KIAA0103 gene product (KIAA0101), mRNA NM 014766 Homo sapiens KIAA0095 gene product (KIAA0095), mRNA NM 014679 Homo sapiens KIAA0092 gene product (KIAA0092), mRNA NM 014679 Homo sapiens KIAA0087 gene product (KIAA0092), mRNA NM 014769 Homo sapiens KIAA0087 gene product (KIAA0094), mRNA NM 014877 Homo sapiens KIAA0087 gene product (KIAA0054), mRNA NM 014716 Homo sapiens KIAA0084 (KIAA0054), mRNA NM 014716 Homo sapiens RIAA0085 (KIAA0054), mRNA NM 014716 Homo sapiens RIAA0088 gene product (KIAA0054), mRNA NM 014716 Homo sapiens RIAA0088 gene product (KIAA0054), mRNA NM 014716 Homo sapiens RIAA0088 gene product (KIAA0054), mRNA NM 014716 Homo sapiens RIAA0088 gene product (KIAA0054), mRNA NM 014716 Homo sapiens RIAA0088 gene product (KIAA0054), mRNA NM 014716 Homo sapiens RIAA0088 gene product (KIAA0054), mRNA
NM 014725 Homo sapiens KIAA0189 gene product (KIAA0189), mRNA NM 014753 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA NM 014791 Homo sapiens likely ortholog of maternal embryonic leucine zipper kinase (KIAA0175), mRNA NM 014746 Homo sapiens KIAA0161 gene product (KIAA0161), mRNA NM 014633 Homo sapiens KIAA0155 gene product (KIAA0155), mRNA NM 014002 Homo sapiens KIK-related kinase epsilon; inducible IkappaB kinase (IKKE), mRNA NM 014847 Homo sapiens KIAA0144 gene product (KIAA0144), mRNA NM 014773 Homo sapiens KIAA0141 gene product (KIAA0141), mRNA NM 014649 Homo sapiens KIAA0138 gene product (KIAA0138), mRNA NM 014792 Homo sapiens KIAA0125 gene product (KIAA0125), mRNA NM 014794 Homo sapiens KIAA0118 protein (KIAA0118), mRNA NM 014740 Homo sapiens KIAA0111 gene product (KIAA0111), mRNA NM 014673 Homo sapiens KIAA0103 gene product (KIAA0103), mRNA NM 01469 Homo sapiens KIAA0109 gene product (KIAA0101), mRNA NM 01469 Homo sapiens KIAA0095 gene product (KIAA0095), mRNA NM 014679 Homo sapiens KIAA0092 gene product (KIAA0095), mRNA NM 014679 Homo sapiens KIAA0092 gene product (KIAA0095), mRNA NM 014769 Homo sapiens KIAA0092 gene product (KIAA0095), mRNA NM 014769 Homo sapiens KIAA0087 gene product (KIAA0095), mRNA NM 014769 Homo sapiens KIAA0087 gene product (KIAA0087), mRNA NM 014716 Homo sapiens R3H domain (binds single-stranded nucleic acids) containing (R3HDM), mRNA
NM_014753 Homo sapiens KIAA0187 gene product (KIAA0187), mRNA NM_014791 Homo sapiens likely ortholog of maternal embryonic leucine zipper kinase (KIAA0175), mRNA NM_014746 Homo sapiens KIAA0161 gene product (KIAA0161), mRNA NM_014633 Homo sapiens KIAA0155 gene product (KIAA0155), mRNA NM_014002 Homo sapiens KIAA0155 gene product (KIAA0155), mRNA NM_014002 Homo sapiens KIAA0144 gene product (KIAA0144), mRNA NM_014847 Homo sapiens KIAA0141 gene product (KIAA0144), mRNA NM_014773 Homo sapiens KIAA0141 gene product (KIAA0141), mRNA NM_014649 Homo sapiens KIAA0138 gene product (KIAA0138), mRNA NM_014792 Homo sapiens KIAA0125 gene product (KIAA0125), mRNA NM_014790 Homo sapiens KIAA0111 gene product (KIAA0111), mRNA NM_014740 Homo sapiens KIAA0111 gene product (KIAA0111), mRNA NM_014673 Homo sapiens KIAA0103 gene product (KIAA0103), mRNA NM_014669 Homo sapiens KIAA0095 gene product (KIAA0101), mRNA NM_014669 Homo sapiens KIAA0095 gene product (KIAA0092), mRNA NM_014679 Homo sapiens KIAA0092 gene product (KIAA0092), mRNA NM_014769 Homo sapiens KIAA0087 gene product (KIAA0087), mRNA NM_014769 Homo sapiens KIAA0087 gene product (KIAA0087), mRNA NM_014769 Homo sapiens RIAA0087 gene product (KIAA0087), mRNA NM_014760 Homo sapiens RIAA0087 gene product (KIAA0087), mRNA NM_014761 Homo sapiens R3H domain (binds single-stranded nucleic acids) containing (R3HDM), mRNA
NM_014791 Homo sapiens likely ortholog of maternal embryonic leucine zipper kinase (KIAA0175), mRNA  NM_014634 Homo sapiens KIAA0161 gene product (KIAA0161), mRNA  NM_014002 Homo sapiens KIAA0155 gene product (KIAA0155), mRNA  NM_014847 Homo sapiens KIAA0144 gene product (KIAA0144), mRNA  NM_014773 Homo sapiens KIAA0141 gene product (KIAA0144), mRNA  NM_014649 Homo sapiens KIAA0138 gene product (KIAA0138), mRNA  NM_014792 Homo sapiens KIAA0125 gene product (KIAA0138), mRNA  NM_014790 Homo sapiens KIAA0118 protein (KIAA0118), mRNA  NM_014740 Homo sapiens KIAA0111 gene product (KIAA0111), mRNA  NM_014736 Homo sapiens KIAA0103 gene product (KIAA0103), mRNA  NM_014736 Homo sapiens KIAA0101 gene product (KIAA0101), mRNA  NM_014699 Homo sapiens KIAA0095 gene product (KIAA0101), mRNA  NM_014736 Homo sapiens KIAA0095 gene product (KIAA0095), mRNA  NM_01469 Homo sapiens KIAA0095 gene product (KIAA0092), mRNA  NM_014769 Homo sapiens KIAA0087 gene product (KIAA0087), mRNA  NM_014769 Homo sapiens KIAA0087 gene product (KIAA0087), mRNA  NM_014716 Homo sapiens Centaurin, beta 1 (CENTB1), mRNA  NM_014716 Homo sapiens R3H domain (binds single-stranded nucleic acids) containing (R3HDM), mRNA
(KIAA0175), mRNA  NM_014746 Homo sapiens KIAA0161 gene product (KIAA0161), mRNA  NM_014633 Homo sapiens KIAA0155 gene product (KIAA0155), mRNA  NM_014002 Homo sapiens IKK-related kinase epsilon; inducible IkappaB kinase (IKKE), mRNA  NM_014847 Homo sapiens KIAA0144 gene product (KIAA0144), mRNA  NM_01473 Homo sapiens KIAA0141 gene product (KIAA0141), mRNA  NM_014649 Homo sapiens KIAA0138 gene product (KIAA0138), mRNA  NM_014792 Homo sapiens KIAA0125 gene product (KIAA0125), mRNA  NM_014999 Homo sapiens KIAA0118 protein (KIAA0118), mRNA  NM_014740 Homo sapiens KIAA0111 gene product (KIAA0111), mRNA  NM_014673 Homo sapiens KIAA0103 gene product (KIAA0103), mRNA  NM_014736 Homo sapiens KIAA0101 gene product (KIAA0101), mRNA  NM_014669 Homo sapiens KIAA0095 gene product (KIAA0095), mRNA  NM_014679 Homo sapiens KIAA0092 gene product (KIAA0092), mRNA  NM_01476 Homo sapiens KIAA0087 gene product (KIAA0087), mRNA  NM_01476 Homo sapiens KIAA0087 gene product (KIAA0087), mRNA  NM_01476 Homo sapiens KIAA0087 gene product (KIAA0087), mRNA  NM_014716 Homo sapiens Centaurin, beta 1 (CENTB1), mRNA  NM_015361 Homo sapiens R3H domain (binds single-stranded nucleic acids) containing (R3HDM), mRNA
NM_014746 Homo sapiens KIAA0161 gene product (KIAA0161), mRNA NM_014633 Homo sapiens KIAA0155 gene product (KIAA0155), mRNA NM_014002 Homo sapiens IKK-related kinase epsilon; inducible IkappaB kinase (IKKE), mRNA NM_014847 Homo sapiens KIAA0144 gene product (KIAA0144), mRNA NM_014773 Homo sapiens KIAA0141 gene product (KIAA0141), mRNA NM_014649 Homo sapiens KIAA0138 gene product (KIAA0138), mRNA NM_014792 Homo sapiens KIAA0125 gene product (KIAA0125), mRNA NM_014999 Homo sapiens KIAA0118 protein (KIAA0118), mRNA NM_014740 Homo sapiens KIAA0111 gene product (KIAA0111), mRNA NM_014673 Homo sapiens KIAA0103 gene product (KIAA0103), mRNA NM_014736 Homo sapiens KIAA0101 gene product (KIAA0101), mRNA NM_014669 Homo sapiens KIAA0095 gene product (KIAA0095), mRNA NM_014679 Homo sapiens KIAA0092 gene product (KIAA0092), mRNA NM_014769 Homo sapiens KIAA0087 gene product (KIAA0087), mRNA NM_014769 Homo sapiens KIAA0087 gene product (KIAA0054), mRNA NM_014716 Homo sapiens centaurin, beta 1 (CENTB1), mRNA NM_014716 Homo sapiens R3H domain (binds single-stranded nucleic acids) containing (R3HDM), mRNA
NM 014633 Homo sapiens KIAA0155 gene product (KIAA0155), mRNA  NM 014002 Homo sapiens IKK-related kinase epsilon; inducible IkappaB kinase (IKKE), mRNA  NM 014847 Homo sapiens KIAA0144 gene product (KIAA0144), mRNA  NM 014773 Homo sapiens KIAA0141 gene product (KIAA0141), mRNA  NM 014649 Homo sapiens KIAA0138 gene product (KIAA0138), mRNA  NM 014792 Homo sapiens KIAA0125 gene product (KIAA0125), mRNA  NM 014999 Homo sapiens KIAA0118 protein (KIAA0118), mRNA  NM 014740 Homo sapiens KIAA0111 gene product (KIAA0111), mRNA  NM 014673 Homo sapiens KIAA0103 gene product (KIAA0103), mRNA  NM 014769 Homo sapiens KIAA0101 gene product (KIAA0101), mRNA  NM 014669 Homo sapiens KIAA0095 gene product (KIAA0095), mRNA  NM 014679 Homo sapiens KIAA0092 gene product (KIAA0087), mRNA  NM 014769 Homo sapiens KIAA0087 gene product (KIAA0087), mRNA  NM 014877 Homo sapiens KIAA0087 gene product (KIAA0054), mRNA  NM 014716 Homo sapiens R3H domain (binds single-stranded nucleic acids) containing (R3HDM), mRNA
NM_014847 Homo sapiens KIAA0144 gene product (KIAA0144), mRNA  NM_014847 Homo sapiens KIAA0141 gene product (KIAA0144), mRNA  NM_014773 Homo sapiens KIAA0141 gene product (KIAA0141), mRNA  NM_014649 Homo sapiens KIAA0138 gene product (KIAA0138), mRNA  NM_014792 Homo sapiens KIAA0125 gene product (KIAA0125), mRNA  NM_014999 Homo sapiens KIAA0118 protein (KIAA0118), mRNA  NM_014740 Homo sapiens KIAA0111 gene product (KIAA0111), mRNA  NM_014673 Homo sapiens KIAA0103 gene product (KIAA0103), mRNA  NM_014736 Homo sapiens KIAA0101 gene product (KIAA0101), mRNA  NM_014669 Homo sapiens KIAA0095 gene product (KIAA0095), mRNA  NM_014679 Homo sapiens KIAA0092 gene product (KIAA0092), mRNA  NM_014769 Homo sapiens KIAA0087 gene product (KIAA0087), mRNA  NM_014877 Homo sapiens kIAA0087 gene product (KIAA0054), mRNA  NM_014716 Homo sapiens centaurin, beta 1 (CENTB1), mRNA  NM_015361 Homo sapiens R3H domain (binds single-stranded nucleic acids) containing (R3HDM), mRNA
mRNA  NM 014847 Homo sapiens KIAA0144 gene product (KIAA0144), mRNA  NM 014773 Homo sapiens KIAA0141 gene product (KIAA0141), mRNA  NM 014649 Homo sapiens KIAA0138 gene product (KIAA0138), mRNA  NM 014792 Homo sapiens KIAA0125 gene product (KIAA0125), mRNA  NM 014999 Homo sapiens KIAA0118 protein (KIAA0118), mRNA  NM 014740 Homo sapiens KIAA0111 gene product (KIAA0111), mRNA  NM 014673 Homo sapiens KIAA0103 gene product (KIAA0103), mRNA  NM 014736 Homo sapiens KIAA0101 gene product (KIAA0101), mRNA  NM 014669 Homo sapiens KIAA0095 gene product (KIAA0095), mRNA  NM 014679 Homo sapiens KIAA0092 gene product (KIAA0092), mRNA  NM 014769 Homo sapiens KIAA0087 gene product (KIAA0087), mRNA  NM 014877 Homo sapiens kIAA0087 gene product (KIAA0054), mRNA  NM 014716 Homo sapiens centaurin, beta 1 (CENTB1), mRNA  NM 015361 Homo sapiens R3H domain (binds single-stranded nucleic acids) containing (R3HDM), mRNA
NM 014847 Homo sapiens KIAA0144 gene product (KIAA0144), mRNA NM 014773 Homo sapiens KIAA0141 gene product (KIAA0141), mRNA NM 014649 Homo sapiens KIAA0138 gene product (KIAA0138), mRNA NM 014792 Homo sapiens KIAA0125 gene product (KIAA0125), mRNA NM 014999 Homo sapiens KIAA0118 protein (KIAA0118), mRNA NM 014740 Homo sapiens KIAA0111 gene product (KIAA0111), mRNA NM 014673 Homo sapiens KIAA0103 gene product (KIAA0103), mRNA NM 014736 Homo sapiens KIAA0101 gene product (KIAA0101), mRNA NM 014669 Homo sapiens KIAA0095 gene product (KIAA0095), mRNA NM 014679 Homo sapiens KIAA0092 gene product (KIAA0092), mRNA NM 014769 Homo sapiens KIAA0087 gene product (KIAA0087), mRNA NM 014877 Homo sapiens kIAA0087 gene product (KIAA0054), mRNA NM 014716 Homo sapiens centaurin, beta 1 (CENTB1), mRNA NM 015361 Homo sapiens R3H domain (binds single-stranded nucleic acids) containing (R3HDM), mRNA
NM 014679 Homo sapiens KIAA0141 gene product (KIAA0141), mRNA NM 01469 Homo sapiens KIAA0138 gene product (KIAA0138), mRNA NM 014792 Homo sapiens KIAA0125 gene product (KIAA0125), mRNA NM 014999 Homo sapiens KIAA0118 protein (KIAA0118), mRNA NM 014740 Homo sapiens KIAA0111 gene product (KIAA0111), mRNA NM 014673 Homo sapiens KIAA0103 gene product (KIAA0103), mRNA NM 014736 Homo sapiens KIAA0101 gene product (KIAA0101), mRNA NM 014669 Homo sapiens KIAA0095 gene product (KIAA0095), mRNA NM 014679 Homo sapiens KIAA0092 gene product (KIAA0092), mRNA NM 014769 Homo sapiens KIAA0087 gene product (KIAA0087), mRNA NM 014877 Homo sapiens kIAA0087 gene product (KIAA0054), mRNA NM 014716 Homo sapiens centaurin, beta 1 (CENTB1), mRNA NM 015361 Homo sapiens R3H domain (binds single-stranded nucleic acids) containing (R3HDM), mRNA
NM 014649 Homo sapiens KIAA0138 gene product (KIAA0138), mRNA NM 014792 Homo sapiens KIAA0125 gene product (KIAA0125), mRNA NM 014999 Homo sapiens KIAA0118 protein (KIAA0118), mRNA NM 014740 Homo sapiens KIAA0111 gene product (KIAA0111), mRNA NM 014673 Homo sapiens KIAA0103 gene product (KIAA0103), mRNA NM 014736 Homo sapiens KIAA0101 gene product (KIAA0101), mRNA NM 014669 Homo sapiens KIAA0095 gene product (KIAA0095), mRNA NM 014679 Homo sapiens KIAA0092 gene product (KIAA0092), mRNA NM 014769 Homo sapiens KIAA0087 gene product (KIAA0087), mRNA NM 014877 Homo sapiens helicase KIAA0054 (KIAA0054), mRNA NM 014716 Homo sapiens centaurin, beta 1 (CENTB1), mRNA NM 015361 Homo sapiens R3H domain (binds single-stranded nucleic acids) containing (R3HDM), mRNA
NM_014792 Homo sapiens KIAA0125 gene product (KIAA0125), mRNA  NM_014999 Homo sapiens KIAA0118 protein (KIAA0118), mRNA  NM_014740 Homo sapiens KIAA0111 gene product (KIAA0111), mRNA  NM_014673 Homo sapiens KIAA0103 gene product (KIAA0103), mRNA  NM_014736 Homo sapiens KIAA0101 gene product (KIAA0101), mRNA  NM_014669 Homo sapiens KIAA0095 gene product (KIAA0095), mRNA  NM_014679 Homo sapiens KIAA0092 gene product (KIAA0092), mRNA  NM_014769 Homo sapiens KIAA0087 gene product (KIAA0087), mRNA  NM_014877 Homo sapiens kIAA0087 gene product (KIAA0054), mRNA  NM_014716 Homo sapiens centaurin, beta 1 (CENTB1), mRNA  NM_015361 Homo sapiens R3H domain (binds single-stranded nucleic acids) containing (R3HDM), mRNA
NM 014999 Homo sapiens KIAA0118 protein (KIAA0118), mRNA  NM_014740 Homo sapiens KIAA0111 gene product (KIAA0111), mRNA  NM_014673 Homo sapiens KIAA0103 gene product (KIAA0103), mRNA  NM_014736 Homo sapiens KIAA0101 gene product (KIAA0101), mRNA  NM_014669 Homo sapiens KIAA0095 gene product (KIAA0095), mRNA  NM_014679 Homo sapiens KIAA0092 gene product (KIAA0092), mRNA  NM_014769 Homo sapiens KIAA0087 gene product (KIAA0087), mRNA  NM_014877 Homo sapiens helicase KIAA0054 (KIAA0054), mRNA  NM_014716 Homo sapiens centaurin, beta 1 (CENTB1), mRNA  NM_015361 Homo sapiens R3H domain (binds single-stranded nucleic acids) containing (R3HDM), mRNA
NM 014740 Homo sapiens KIAA0111 gene product (KIAA0111), mRNA  NM 014673 Homo sapiens KIAA0103 gene product (KIAA0103), mRNA  NM 014736 Homo sapiens KIAA0101 gene product (KIAA0101), mRNA  NM 014669 Homo sapiens KIAA0095 gene product (KIAA0095), mRNA  NM 014679 Homo sapiens KIAA0092 gene product (KIAA0092), mRNA  NM 014769 Homo sapiens KIAA0087 gene product (KIAA0087), mRNA  NM 014877 Homo sapiens kIAA0087 (KIAA0054), mRNA  NM 014716 Homo sapiens centaurin, beta 1 (CENTB1), mRNA  NM 015361 Homo sapiens R3H domain (binds single-stranded nucleic acids) containing (R3HDM), mRNA
NM 014673 Homo sapiens KIAA0103 gene product (KIAA0103), mRNA  NM 014736 Homo sapiens KIAA0101 gene product (KIAA0101), mRNA  NM 014669 Homo sapiens KIAA0095 gene product (KIAA0095), mRNA  NM 014679 Homo sapiens KIAA0092 gene product (KIAA0092), mRNA  NM 014769 Homo sapiens KIAA0087 gene product (KIAA0087), mRNA  NM 014877 Homo sapiens helicase KIAA0054 (KIAA0054), mRNA  NM 014716 Homo sapiens centaurin, beta 1 (CENTB1), mRNA  NM 015361 Homo sapiens R3H domain (binds single-stranded nucleic acids) containing (R3HDM), mRNA
NM 014736 Homo sapiens KIAA0101 gene product (KIAA0101), mRNA NM 014669 Homo sapiens KIAA0095 gene product (KIAA0095), mRNA NM 014679 Homo sapiens KIAA0092 gene product (KIAA0092), mRNA NM 014769 Homo sapiens KIAA0087 gene product (KIAA0087), mRNA NM 014877 Homo sapiens helicase KIAA0054 (KIAA0054), mRNA NM 014716 Homo sapiens centaurin, beta 1 (CENTB1), mRNA NM 015361 Homo sapiens R3H domain (binds single-stranded nucleic acids) containing (R3HDM), mRNA
NM_014669 Homo sapiens KIAA0095 gene product (KIAA0095), mRNA NM_014679 Homo sapiens KIAA0092 gene product (KIAA0092), mRNA NM_014769 Homo sapiens KIAA0087 gene product (KIAA0087), mRNA NM_014877 Homo sapiens helicase KIAA0054 (KIAA0054), mRNA NM_014716 Homo sapiens centaurin, beta 1 (CENTB1), mRNA NM_015361 Homo sapiens R3H domain (binds single-stranded nucleic acids) containing (R3HDM), mRNA
NM_014679 Homo sapiens KIAA0092 gene product (KIAA0092), mRNA NM_014769 Homo sapiens KIAA0087 gene product (KIAA0087), mRNA NM_014877 Homo sapiens helicase KIAA0054 (KIAA0054), mRNA NM_014716 Homo sapiens centaurin, beta 1 (CENTB1), mRNA NM_015361 Homo sapiens R3H domain (binds single-stranded nucleic acids) containing (R3HDM), mRNA
NM_014769 Homo sapiens KIAA0087 gene product (KIAA0087), mRNA  NM_014877 Homo sapiens helicase KIAA0054 (KIAA0054), mRNA  NM_014716 Homo sapiens centaurin, beta 1 (CENTB1), mRNA  NM_015361 Homo sapiens R3H domain (binds single-stranded nucleic acids) containing (R3HDM), mRNA
NM_014877 Homo sapiens helicase KIAA0054 (KIAA0054), mRNA  NM_014716 Homo sapiens centaurin, beta 1 (CENTB1), mRNA  NM_015361 Homo sapiens R3H domain (binds single-stranded nucleic acids) containing (R3HDM), mRNA
NM_014716 Homo sapiens centaurin, beta 1 (CENTB1), mRNA NM_015361 Homo sapiens R3H domain (binds single-stranded nucleic acids) containing (R3HDM), mRNA
NM_015361 Homo sapiens R3H domain (binds single-stranded nucleic acids) containing (R3HDM), mRNA
(R3HDM), mRNA
INIVE U1400U - E HOMO SADIENS KTA AUUZZ VENE DYOGNELIKTA AUUZZE MKINA
NM_014878 Homo sapiens KIAA0020 gene product (KIAA0020), mRNA
NM_014665 Homo sapiens KIAA0014 gene product (KIAA0014), mRNA
NM_014671   Homo sapiens ubiquitin-protein isopeptide ligase (E3) (KIAA0010), mRNA
NM_014637 Homo sapiens KIAA0009 gene product (KIAA0009), mRNA
NM_015384 Homo sapiens IDN3 protein (IDN3), mRNA
NM_014188   Homo sapiens HSPC182 protein (HSPC182), mRNA
NM_014187 Homo sapiens HSPC171 protein (HSPC171), mRNA
NM_014182 Homo sapiens HSPC160 protein (HSPC160), mRNA
NIM DIALTO LITERA
NM_014178 Homo sapiens HSPC156 protein (HSPC156), mRNA NM_014177 Homo sapiens HSPC154 protein (HSPC154), mRNA

ND 6 014176	II III III III III III III III III
NM_014176	Homo sapiens HSPC150 protein similar to ubiquitin-conjugating enzyme
ND 6 014172	(HSPC150), mRNA
NM_014173	Homo sapiens HSPC142 protein (HSPC142), mRNA
NM_014172	Homo sapiens HSPC141 protein (HSPC141), mRNA
NM_014171	Homo sapiens postsynaptic protein CRIPT (CRIPT), mRNA
NM_014169	Homo sapiens HSPC134 protein (HSPC134), mRNA
NM_014168	Homo sapiens HSPC133 protein (HSPC133), mRNA
NM_014167	Homo sapiens HSPC128 protein (HSPC128), mRNA
NM_014165	Homo sapiens HSPC125 protein (HSPC125), mRNA
NM_014163	Homo sapiens HSPC073 protein (HSPC073), mRNA
NM_014162	Homo sapiens HSPC072 protein (HSPC072), mRNA
NM_014159	Homo sapiens Huntingtin interacting protein B (HYPB), mRNA
NM_014158	Homo sapiens HSPC067 protein (HSPC067), mRNA
NM_014157	Homo sapiens HSPC065 protein (HSPC065), mRNA
NM_014152	Homo sapiens HSPC054 protein (HSPC054), mRNA
NM 014151	Homo sapiens HSPC053 protein (HSPC053), mRNA
NM_014148	Homo sapiens HSPC048 protein (HSPC048), mRNA
NM_014147	Homo sapiens HSPC047 protein (HSPC047), mRNA
NM 014041	Homo sapiens signal peptidase 12kDa (SPC12), mRNA
NM 014047	Homo sapiens HSPC023 protein (HSPC023), mRNA
NM 014028	Homo sapiens HSPC019 protein (HSPC019), mRNA
NM 014026	Homo sapiens HSPC015 protein (HSPC015), mRNA
NM 015362	Homo sapiens HSPC002 protein (HSPC002), mRNA
NM 015603	Homo sapiens DKFZP586M1019 protein (DKFZP586M1019), mRNA
NM 015537	Homo sapiens DKFZP586J1624 protein (DKFZP586J1624), mRNA
NM 015584	Homo sapiens DKFZP586F1524 protein (DKFZP586F1524), mRNA
NM 015677	Homo sapiens hypothetical protein (DKFZP586F1318), mRNA
NM 015416	Homo sapiens DKFZP586A011 protein (DKFZP586A011), mRNA
NM 015513	Homo sapiens DKFZP566D213 protein (DKFZP566D213), mRNA
NM 015509	Homo sapiens DKFZP566B183 protein (DKFZP566B183), mRNA
NM_014042	Homo sapiens DKFZP564M082 protein (DKFZP564M082), mRNA
NM_015455	Homo sapiens KIAA1194 protein (KIAA1194), mRNA
NM_015601	Homo sapiens DKFZP564G092 protein (DKFZP564G092), mRNA
NM 014044	Homo sapiens DKFZP564G0222 protein (DKFZP564G0222), mRNA
NM 015658	Homo sapiens DKFZP564C186 protein (DKFZP564C186), mRNA
NM_015654	Homo sapiens DKFZP564C103 protein (DKFZP564C103), mRNA
NM 015535	Homo sapiens DKFZP564A2416 protein (DKFZP564A2416), mRNA
NM 014034	Homo sapiens DKFZP547E2110 protein (DKFZP547E2110), mRNA
NM 015607	Homo sapiens DKFZP547E1010 protein (DKFZP547E1010), mRNA
NM 015594	Homo sapiens DKFZP434O047 protein (DKFZP434O047), mRNA
NM 015492	Homo sapiens DKFZP434H132 protein (DKFZP434H132), mRNA
NM 015515	Homo sapiens type I intermediate filament cytokeratin (HAIK1), mRNA
NM 014064	Homo sapiens AD-003 protein (AD-003), mRNA
NM 014517	Homo sapiens upstream binding protein 1 (LBP-1a) (UBP1), mRNA
NM 014294	Homo sapiens translocating chain-associating membrane protein (TRAM),
	mRNA
NM_014305	Homo sapiens dTDP-D-glucose 4,6-dehydratase (TDPGD), mRNA
NM 014300	Homo sapiens signal peptidase complex (18kD) (SPC18), mRNA
NM 014419	Homo sapiens soggy-1 gene (DKKL1-pending), mRNA
NM 014445	Homo sapiens stress-associated endoplasmic reticulum protein 1; ribosome
_===	associated membrane protein 4 (SERP1), mRNA
NM 014329	Homo sapiens autoantigen (RCD-8), mRNA

NM_014504	Homo sapiens putative Rab5 GDP/GTP exchange factor homologue (RABEX5), mRNA
NM 014589	Homo sapiens phospholipase A2, group IIE (PLA2G2E), mRNA
NM 014471	Homo sapiens serine protease inhibitor, Kazal type 4 (SPINK4), mRNA
NM 014891	Homo sapiens PDGFA associated protein 1 (PDAP1), mRNA
NM 014308	Homo sapiens phosphoinositide-3-kinase, regulatory subunit, polypeptide p101
_	(P101-PI3K), mRNA
NM 014359	Homo sapiens opticin (OPTC), mRNA
NM 014515	Homo sapiens CCR4-NOT transcription complex, subunit 2 (CNOT2), mRNA
NM 014360	Homo sapiens NK-2 (Drosophila) homolog 8 (NKX2.8), mRNA
NM 014371	Homo sapiens neighbor of A-kinase anchoring protein 95 (NAKAP95), mRNA
NM 014342	Homo sapiens mitochondrial carrier homolog 2 (MTCH2), nuclear gene
_	encoding mitochondrial protein, mRNA
NM 015716	Homo sapiens Misshapen/NIK-related kinase (MINK), mRNA
NM 014358	Homo sapiens C-type (calcium dependent, carbohydrate-recognition domain)
-	lectin, superfamily member 9 (CLECSF9), mRNA
NM_014552	Homo sapiens LBP protein 32 (LBP-32), mRNA
NM_014247	Homo sapiens PDZ domain containing guanine nucleotide exchange
_	factor(GEF)1 (PDZ-GEF1), mRNA
NM_014267	Homo sapiens small acidic protein (IMAGE145052), mRNA
NM_014597	Homo sapiens acidic 82 kDa protein mRNA (HSU15552), mRNA
NM_014254	Homo sapiens transmembrane protein 5 (TMEM5), mRNA
NM 014362	Homo sapiens 3-hydroxyisobutyryl-Coenzyme A hydrolase (HIBCH), mRNA
NM_014365	Homo sapiens protein kinase H11 (H11), mRNA
NM_014584	Homo sapiens ERO1-like (S. cerevisiae) (ERO1L), mRNA
NM_014367	Homo sapiens hypothetical protein, estradiol-induced (E2IG5), mRNA
NM_014366	Homo sapiens putative nucleotide binding protein, estradiol-induced (E2IG3), mRNA
NM_014380	Homo sapiens nerve growth factor receptor (TNFRSF16) associated protein 1 (NGFRAP1), mRNA
NM 014890	Homo sapiens downregulated in ovarian cancer 1 (DOC1), mRNA
NM_014595	Homo sapiens 5' nucleotidase, deoxy (pyrimidine), cytosolic type C (NT5C), mRNA
NM_014316	Homo sapiens calcium-regulated heat-stable protein (24kD) (CRHSP-24), mRNA
NM 014430	Homo sapiens cell death-inducing DFFA-like effector b (CIDEB), mRNA
NM_014400	Homo sapiens GPI-anchored metastasis-associated protein homolog (C4.4A), mRNA
NM_014408	Homo sapiens similar to yeast BET3 (S. cerevisiae) (BET3), mRNA
NM_014374	Homo sapiens replication initiation region protein (60kD) (RIP60), mRNA
NM_013943	Homo sapiens chloride intracellular channel 4 (CLIC4), mRNA
NM_013433	Homo sapiens karyopherin beta 2b, transportin (TRN2), mRNA
NM_013435	Homo sapiens retinal homeobox protein (RX), mRNA
NM_013377	Homo sapiens hypothetical protein (DKFZp434B0417), mRNA
NM_012297	Homo sapiens Ras-GTPase activating protein SH3 domain-binding protein 2
	(KIAA0660), mRNA
NM_013286	Homo sapiens chromosome 3p21.1 gene sequence (HUMAGCGB), mRNA
NM_012472	Homo sapiens testis specific leucine rich repeat protein (TSLRP), mRNA
NM_012119	Homo sapiens cell cycle related kinase (CCRK), mRNA
NM_013266	Homo sapiens alpha-catenin-like protein (VR22), mRNA
NM_013346	Homo sapiens sorting nexin 12 (SNX12), mRNA
NM 013322	Homo sapiens sorting nexin 10 (SNX10), mRNA

376 010	
NM_013400	Homo sapiens replication initiation region protein (60kD) (RIP60), mRNA
NM_013355	Homo sapiens protein kinase PKNbeta (pknbeta), mRNA
NM_013240	Homo sapiens putative N6-DNA-methyltransferase (N6AMT1) mRNA
NM_013364	Homo sapiens paraneoplastic cancer-testis-brain antigen (MAS) mPNA
NM_013275	Homo sapiens nasopharyngeal carcinoma susceptibility protein (I.Z16), mRNA
NM_013312	Homo sapiens nook2 protein (HOOK2), mRNA
NM_013332	Homo sapiens hypoxia-inducible protein 2 (HIG2), mRNA
NM_013308	Homo sapiens platelet activating receptor homolog (H963) mPNA
NM 013394	Homo sapiens acid fibroblast growth factor-like protein (GLIO703) mPNA
NM_013329	nomo sapiens chromosome 21 open reading frame 66 (C21orf66) mDNA
NM_013333	Homo sapiens EH domain-binding mitotic phosphoprotein (EPSIN) mRNA
NM_013395	Holio sapiens proteinx0008 (AD013), mRNA
NM_012463	Homo sapiens TJ6 protein (TJ6), mRNA
NM_012461	Homo sapiens TERF1 (TRF1)-interacting nuclear factor 2 (TINF2), mRNA
NM_012245	110110 sapiens SK1-interacting protein (SNW1) mRNA
NM_012437	Homo sapiens SNARE associated protein snapin (SNAPAP) mRNA
NM_012433	Homo sapiens splicing factor 3b, subunit 1, 155kD (SF3R1) mPNA
NM_012431	Homo sapiens sema domain, immunoglobulin domain (Ig) short basis domain
277.6.01000.1	1 secreted, (semaphorm) 3E (SEMA3E), mRNA
NM_012234	Homo sapiens RING1 and YY1 binding protein (RVRP) mPNA
NM_012420	Homo sapiens retinoic acid- and interferon-inducible protein (58kD) (R158)
ND 6 010415	IIIKINA
NM_012417	Homo sapiens retinal degeneration B beta (RDGBB), mRNA
NM_012229	Homo sapiens 5'-nucleotidase (purine), cytosolic type R (NTSP), mp.NA
NM_012390	Homo sapiens protein homologous to salivary proline-rich protein P.B. (PDD)
ND4 012246	IIICIVA
NM_012346	Homo sapiens nucleoporin 62kD (NUP62), mRNA
NM_012339	Homo sapiens transmembrane 4 superfamily member (tetraspan NET-7) (NET-7) mPNA
NM 012338	1 /), nuclea
1111_012556	Homo sapiens transmembrane 4 superfamily member (tetraspan NET-2) (NET-2), mRNA
NM_012332	
NM_012327	Homo sapiens Mitochondrial Acyl-CoA Thioesterase (MT-ACT48), mRNA
NM_012321	Homo sapiens phosphatidylinositol glycan class N (PIGN) mPNA
NM_012294	Homo sapiens U6 snRNA-associated Sm-like protein (LSM4), mRNA
11112_012254	Homo sapiens guanine nucleotide exchange factor for Rap1; M-Ras-regulated GEF (KIAA0277), mRNA
NM 012289	Homo senions Volab III. FOUR
NM_012285	Homo sapiens Kelch-like ECH-associated protein 1 (KIAA0132), mRNA
	Homo sapiens potassium voltage-gated channel, subfamily H (eag-related), member 4 (KCNH4), mRNA
NM_012267	Homo sapiens hsp70-interacting protein (HSPBP1), mRNA
NM_012266	Homo saniens Dna I (Hendo) home les and Carlo Saniens Dna I (Hendo) home les a
	Homo sapiens DnaJ (Hsp40) homolog, subfamily B, member 5 (DNAJB5), mRNA
NM_012260	
NM 012204	Homo sapiens 2-hydroxyphytanoyl-CoA lyase (HPCL2), mRNA  Homo sapiens general transcription factor HIC - 1
	Homo sapiens general transcription factor IIIC, polypeptide 4 (90kD) (GTF3C4), mRNA
NM_012086	Homo sapiens general transcription factor IIIC, polypeptide 3 (102kD)
	(GTF3C3), mRNA
NM_012155	Homo sapiens microtubule-associated protein like echinoderm EMAP (EMAP-
	2), mRNA
NM_012123	Homo sapiens CGI-02 protein (CGI-02), mRNA
NM_012097	Homo sapiens ADP-ribosylation factor-like 5 (ARL5), mRNA
NM_005028	Homo sapiens phosphatidylinositol-4-phosphate 5-kinase, type II, alpha
	phosphatidy intostror-4-phosphate 5-kinase, type II, alpha

	(PIP5K2A), mRNA
NM 006869	Homo sapiens centaurin, alpha 1 (CENTA1), mRNA
NM 007362	Homo sapiens nuclear cap binding protein subunit 2, 20kD (NCBP2), mRNA
NM 007358	Homo sapiens putative DNA binding protein (M96), mRNA
NM 007338	Homo sapiens transcription termination factor, RNA polymerase I (TTF1).
11111_007544	mRNA
NM 007369	Homo sapiens G-protein coupled receptor (RE2), mRNA
NM 005176	Homo sapiens ATP synthase, H+ transporting, mitochondrial F0 complex,
<del></del>	subunit c (subunit 9), isoform 2 (ATP5G2), mRNA
NM_007347	Homo sapiens adaptor-related protein complex 4, epsilon 1 subunit (AP4E1),
_	mRNA
NM_002673	Homo sapiens plexin B1 (PLXNB1), mRNA
NM_007034	Homo sapiens DnaJ (Hsp40) homolog, subfamily B, member 4 (DNAJB4),
	mRNA
NM_004547	Homo sapiens NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 4 (15kD,
	B15) (NDUFB4), mRNA
NM_007180	Homo sapiens trehalase (brush-border membrane glycoprotein) (TREH), mRNA
NM_007115	Homo sapiens tumor necrosis factor, alpha-induced protein 6 (TNFAIP6),
	mRNA
NM_007217	Homo sapiens programmed cell death 10 (PDCD10), mRNA
NM_007269	Homo sapiens syntaxin binding protein 3 (STXBP3), mRNA
NM_007107	Homo sapiens signal sequence receptor, gamma (translocon-associated protein
	gamma) (SSR3), mRNA
NM_007282	Homo sapiens ring finger protein 13 (RNF13), mRNA
NM_007265	Homo sapiens suppressor of S. cerevisiae gcr2 (HSGT1), mRNA
NM_007223	Homo sapiens putative G protein coupled receptor (GPR), mRNA
NM_007192	Homo sapiens chromatin-specific transcription elongation factor, 140 kDa subunit (FACTP140), mRNA
NM_007263	Homo sapiens coatomer protein complex, subunit epsilon (COPE), mRNA
NM_007005	Homo sapiens BCE-1 protein (BCE-1), mRNA
NM_007019	Homo sapiens ubiquitin-conjugating enzyme E2C (UBE2C), mRNA
NM_007064	Homo sapiens serine/threonine kinase with Dbl- and pleckstrin homology
	domains (TRAD), mRNA
NM_007062	Homo sapiens nuclear phosphoprotein similar to S. cerevisiae PWP1 (PWP1), mRNA
NM_007080	Homo sapiens Sm protein F (LSM6), mRNA
NM_007072	Homo sapiens HERV-H LTR-associating 2 (HHLA2), mRNA
NM_007077	Homo sapiens adaptor-related protein complex 4, sigma 1 subunit (AP4S1),
NIM 006751	mRNA
NM_006751	Homo sapiens sperm specific antigen 2 (SSFA2), mRNA
NM_006748	Homo sapiens Src-like-adaptor (SLA), mRNA
NM 006851	Homo sapiens glioma pathogenesis-related protein (RTVP1), mRNA
NM_006815	Homo sapiens coated vesicle membrane protein (RNP24), mRNA
NM_006741	Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 1A (PPP1R1A), mRNA
NM_006823	Homo sapiens protein kinase (cAMP-dependent, catalytic) inhibitor alpha
	(PKIA), mRNA
NM_006825	Homo sapiens cytoskeleton-associated protein 4 (CKAP4), mRNA
NM_006833	Homo sapiens COP9 subunit 6 (MOV34 homolog, 34 kD) (MOV34-34KD),
	mRNA
NM_006838	Homo sapiens methionyl aminopeptidase 2 (METAP2), mRNA
	Homo sapiens vesicle-associated membrane protein 5 (myobrevin) (VAMP5),

orotein
orotein
rotein
rotein
OP4),
JP4),
),
),
2)
-),
<u> </u>
,
35),
35),
NA
12.2
(v-
F
C1A),
mplex
mpiox
PEF2),
(v- F

	mRNA
NM_006230	Homo sapiens polymerase (DNA directed), delta 2, regulatory subunit (50kD)
	(POLD2), mRNA
NM_006156	Homo sapiens neural precursor cell expressed, developmentally down-regulated
	8 (NEDD8), mRNA
NM_006369	Homo sapiens MUF1 protein (MUF1), mRNA
NM_006441	Homo sapiens 5,10-methenyltetrahydrofolate synthetase (5-
	formyltetrahydrofolate cyclo-ligase) (MTHFS), mRNA
NM_006309	Homo sapiens leucine rich repeat (in FLII) interacting protein 2 (LRRFIP2), mRNA
NM_006330	Homo sapiens lysophospholipase I (LYPLA1), mRNA
NM_006344	Homo sapiens macrophage lectin 2 (calcium dependent) (HML2), mRNA
NM_006395	Homo sapiens ubiquitin activating enzyme E1-like protein (GSA7), mRNA
NM_006322	Homo sapiens spindle pole body protein (GCP3), mRNA
NM_006141	Homo sapiens dynein, cytoplasmic, light intermediate polypeptide 2 (DNCLI2), mRNA
NM_006416	Homo sapiens solute carrier family 35 (CMP-sialic acid transporter), member 1 (SLC35A1), mRNA
NM 006349	Homo sapiens putative cyclin G1 interacting protein (CG1I), mRNA
NM 006429	Homo sapiens chaperonin containing TCP1, subunit 7 (eta) (CCT7), mRNA
NM 006430	Homo sapiens chaperonin containing TCP1, subunit 4 (delta) (CCT4), mRNA
NM 006431	Homo sapiens chaperonin containing TCP1, subunit 2 (beta) (CCT2), mRNA
NM 002810	Homo sapiens proteasome (prosome, macropain) 26S subunit, non-ATPase, 4
	(PSMD4), mRNA
NM_006002	Homo sapiens ubiquitin carboxyl-terminal esterase L3 (ubiquitin thiolesterase) (UCHL3), mRNA
NM 006068	Homo sapiens toll-like receptor 6 (TLR6), mRNA
NM 006100	Homo sapiens alpha2,3-sialyltransferase (ST3GALVI), mRNA
NM 006061	
NM 006063	Homo sapiens specific granule protein (28 kDa) (SGP28), mRNA
NM 006076	Homo sapiens sarcomeric muscle protein (SARCOSIN), mRNA  Homo sapiens Boy/Roy activation demain binding matrix alabel (RAP, R)
1414_000076	Homo sapiens Rev/Rex activation domain binding protein-related (RAB-R), mRNA
NM_006034	Homo sapiens p53-induced protein (PIG11), mRNA
NM_006039	Homo sapiens endocytic receptor (macrophage mannose receptor family) (KIAA0709), mRNA
NM_006018	Homo sapiens putative chemokine receptor; GTP-binding protein (HM74), mRNA
NM_006101	Homo sapiens highly expressed in cancer, rich in leucine heptad repeats (HEC), mRNA
NM_006098	Homo sapiens guanine nucleotide binding protein (G protein), beta polypeptide 2-like 1 (GNB2L1), mRNA
NM 005895	Homo sapiens golgi autoantigen, golgin subfamily a, 3 (GOLGA3), mRNA
NM 006023	Homo sapiens D123 gene product (D123), mRNA
NM 006090	Homo sapiens choline/ethanolaminephosphotransferase (CEPT1), mRNA
NM 005822	Homo sapiens Down syndrome critical region gene 1-like 1 (DSCR1L1), mRNA
NM 005827	Homo sapiens UDP-galactose transporter related (UGTREL1), mRNA
NM 005725	Homo sapiens tetraspan 2 (TSPAN-2), mRNA
NM 005879	Homo sapiens TRAF interacting protein (TRIP), mRNA
NM 005816	Homo sapiens T cell activation, increased late expression (TACTILE), mRNA
NM_005843	Homo sapiens signal transducing adaptor molecule (SH3 domain and ITAM
ND ( 005626	motif) 2 (STAM2), mRNA
NM_005636	Homo sapiens synovial sarcoma, X breakpoint 4 (SSX4), mRNA

NM_005775	Homo sapiens vinexin beta (SH3-containing adaptor molecule-1) (SCAM-1), mRNA
NM 005785	Homo sapiens hypothetical SBBI03 protein (SBB103), mRNA
NM 005862	Homo sapiens stromal antigen 1 (STAG1), mRNA
NM 005619	Homo sapiens reticulon 2 (RTN2), mRNA
NM 005615	Homo sapiens ribonuclease, RNase A family, k6 (RNASE6), mRNA
NM 005771	Homo sapiens retinol dehydrogenase homolog (RDHL), mRNA
NM_005833	Homo sapiens Rab9 effector p40 (RAB9P40), mRNA
NM_005687	Homo sapiens phenylalanyl-tRNA synthetase beta-subunit (PheHB), mRNA
NM_005605	Homo sapiens protein phosphatase 3 (formerly 2B), catalytic subunit, gamma isoform (calcineurin A gamma) (PPP3CC), mRNA
NM 005796	Homo sapiens nuclear transport factor 2 (placental protein 15) (PP15), mRNA
NM 005742	Homo sapiens protein disulfide isomerase-related protein (P5), mRNA
NM 005824	Homo sapiens 37 kDa leucine-rich repeat (LRR) protein (P37NB), mRNA
NM 005861	Homo sapiens STIP1 homology and U-Box containing protein 1 (STUB1),
	mRNA
NM_005601	Homo sapiens natural killer cell group 7 sequence (NKG7), mRNA
NM_005831	Homo sapiens nuclear domain 10 protein (NDP52), mRNA
NM_005511	Homo sapiens melan-A (MLANA), mRNA
NM_005575	Homo sapiens leucyl/cystinyl aminopeptidase (LNPEP), mRNA
NM_005794	Homo sapiens short-chain alcohol dehydrogenase family member (HEP27), mRNA
NM_005769	Homo sapiens carbohydrate (N-acetylglucosamine 6-O) sulfotransferase 4 (CHST4), mRNA
NM 005828	Homo sapiens WD-repeat protein (HAN11), mRNA
NM 005804	Homo sapiens nuclear RNA helicase, DECD variant of DEAD box family
	(DDXL), mRNA
NM_005505	Homo sapiens CD36 antigen (collagen type I receptor, thrombospondin receptor)-like 1 (CD36L1), mRNA
NM 005760	Homo sapiens CCAAT-box-binding transcription factor (CBF2), mRNA
NM 005795	Homo sapiens calcitonin receptor-like (CALCRL), mRNA
NM_005720	Homo sapiens actin related protein 2/3 complex, subunit 1B (41 kD) (ARPC1B), mRNA
NM 005876	Homo sapiens nuclear protein, marker for differentiated aortic smooth muscle
_	and down-regulated with vascular injury (APEG1), mRNA
NM_001540	Homo sapiens heat shock 27kD protein 1 (HSPB1), mRNA
NM_005481	Homo sapiens thyroid hormone receptor-associated protein, 95-kD subunit (TRAP95), mRNA
NM 005449	Homo sapiens regulator of Fas-induced apoptosis (TOSO), mRNA
NM 005480	Homo sapiens trophinin associated protein (tastin) (TROAP), mRNA
NM_005419	Homo sapiens signal transducer and activator of transcription 2, 113kD (STAT2), mRNA
NM 005500	Homo sapiens SUMO-1 activating enzyme subunit 1 (SAE1), mRNA
NM 005400	Homo sapiens protein kinase C, epsilon (PRKCE), mRNA
NM 005391	Homo sapiens pyruvate dehydrogenase kinase, isoenzyme 3 (PDK3), mRNA
NM_005494	Homo sapiens DnaJ (Hsp40) homolog, subfamily B, member 6 (DNAJB6), mRNA
NM 005466	Homo sapiens RNA polymerase II transcriptional regulation mediator (Med6, S.
1111_005400	cerevisiae, homolog of) (MED6), mRNA
NM 005310	Homo sapiens growth factor receptor-bound protein 7 (GRB7), mRNA
NM 005497	Homo sapiens gap junction protein, alpha 7, 45kD (connexin 45) (GJA7), mRNA
NM 005175	Homo sapiens ATP synthase, H+ transporting, mitochondrial F0 complex.
	1

	1 1 1 1 ( 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
ND ( 002410	subunit c (subunit 9), isoform 1 (ATP5G1), mRNA
NM_003418	Homo sapiens zinc finger protein 9 (a cellular retroviral nucleic acid binding
1	protein) (ZNF9), mRNA
NM_005151	Homo sapiens ubiquitin specific protease 14 (tRNA-guanine transglycosylase)
	(USP14), mRNA
NM_005119	Homo sapiens thyroid hormone receptor-associated protein, 150 kDa subunit
	(TRAP150), mRNA
NM_005071	Homo sapiens solute carrier family 1 (high affinity aspartate/glutamate
	transporter), member 6 (SLC1A6), mRNA
NM_005047	Homo sapiens proteasome (prosome, macropain) 26S subunit, non-ATPase, 5
	(PSMD5), mRNA
NM_005134	Homo sapiens protein phosphatase 4, regulatory subunit 1 (PPP4R1), mRNA
NM_005033	Homo sapiens polymyositis/scleroderma autoantigen 1 (75kD) (PMSCL1),
	mRNA
NM_005025	Homo sapiens serine (or cysteine) proteinase inhibitor, clade I (neuroserpin),
-	member 1 (SERPINI1), mRNA
NM_005023	Homo sapiens protein geranylgeranyltransferase type I, beta subunit (PGGT1B),
_	mRNA
NM_005020	Homo sapiens phosphodiesterase 1C, calmodulin-dependent (70kD) (PDE1C),
	mRNA
NM_005017	Homo sapiens phosphate cytidylyltransferase 1, choline, alpha isoform
	(PCYT1A), mRNA
NM 005131	Homo sapiens nuclear matrix protein p84 (P84), mRNA
NM 005101	Homo sapiens interferon-stimulated protein, 15 kDa (ISG15), mRNA
NM 005122	Homo sapiens nuclear receptor subfamily 1, group I, member 3 (NR1I3), mRNA
NM 004666	Homo sapiens vanin 1 (VNN1), mRNA
NM 004247	Homo sepiens IIS on PNP specific mentain 1161D (IIS 116KD) PNA
NM 004704	Homo sapiens U5 snRNP-specific protein, 116 kD (U5-116KD), mRNA
NM 004786	Homo sapiens U3 snoRNP-associated 55-kDa protein (U3-55K), mRNA
NM 004257	Homo sapiens thioredoxin-like, 32kD (TXNL), mRNA
NM 004620	Homo sapiens TGF beta receptor associated protein -1 (TRAP-1), mRNA
NM 004604	Homo sapiens TNF receptor-associated factor 6 (TRAF6), mRNA
	Homo sapiens syntaxin 4A (placental) (STX4A), mRNA
NM_004785	Homo sapiens solute carrier family 9 (sodium/hydrogen exchanger), isoform 3
ND 4 004252	regulatory factor 2 (SLC9A3R2), mRNA
NM_004252	Homo sapiens solute carrier family 9 (sodium/hydrogen exchanger), isoform 3
)D/ 004604	regulatory factor 1 (SLC9A3R1), mRNA
NM_004694	Homo sapiens solute carrier family 16 (monocarboxylic acid transporters),
NTM 004505	member 6 (SLC16A6), mRNA
NM_004696	Homo sapiens solute carrier family 16 (monocarboxylic acid transporters),
NTM 004262	member 4 (SLC16A4), mRNA
NM_004263	Homo sapiens sema domain, immunoglobulin domain (Ig), transmembrane
	domain (TM) and short cytoplasmic domain, (semaphorin) 4F (SEMA4F),
3.73.6 00 10 15	mRNA
NM_004868	Homo sapiens glycoprotein, synaptic 2 (GPSN2), mRNA
NM_004844	Homo sapiens SH3-domain binding protein 5 (BTK-associated) (SH3BP5).
	mRNA
NM_004703	Homo sapiens rabaptin-5 (RAB5EP), mRNA
NM_004249	Homo sapiens RAB28, member RAS oncogene family (RAB28), mRNA
NM_004218	Homo sapiens RAB11B, member RAS oncogene family (RAB11B), mRNA
NM_004676	Homo sapiens PTPN13-like, Y-linked (PRY), mRNA
NM_004726	Homo sapiens RALBP1 associated Eps domain containing 2 (REPS2), mRNA
NM_004881	Homo sapiens quinone oxidoreductase homolog (PIG3), mRNA
	, ordeoredance nothering (1 103), mixty

NM_004671	Homo sapiens Protein inhibitor of activated STAT X (PIASX-BETA), mRNA
NM_004565	Homo sapiens peroxisomal biogenesis factor 14 (PEX14), mRNA
NM_004845	Homo sapiens phosphate cytidylyltransferase 1, choline, beta isoform (PCYT1B), mRNA
NM_004563	Homo sapiens phosphoenolpyruvate carboxykinase 2 (mitochondrial) (PCK2), mRNA
NM_004800	Homo sapiens transmembrane 9 superfamily member 2 (TM9SF2), mRNA
NM_004556	Homo sapiens nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor, epsilon (NFKBIE), mRNA
NM 004647	Homo sapiens Neuro-d4 (rat) homolog (NEUD4), mRNA
NM_004546	Homo sapiens NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 2 (8kD, AGGG) (NDUFB2), mRNA
NM_004545	Homo sapiens NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 1 (7kD, MNLL) (NDUFB1), mRNA
NM_004542	Homo sapiens NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 3 (9kD, B9) (NDUFA3), mRNA
NM_004544	Homo sapiens NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 10 (42kD) (NDUFA10), mRNA
NM_004784	Homo sapiens N-deacetylase/N-sulfotransferase (heparan glucosaminyl) 3 (NDST3), mRNA
NM_004901	Homo sapiens lysosomal apyrase-like 1 (LYSAL1), mRNA
NM_004798	Homo sapiens kinesin family member 3B (KIF3B), mRNA
NM_004515	Homo sapiens interleukin enhancer binding factor 2, 45kD (ILF2), mRNA
NM_004838	Homo sapiens Homer, neuronal immediate early gene, 3 (HOMER-3), mRNA
NM_004854	Homo sapiens HNK-1 sulfotransferase (HNK-1ST), mRNA
NM_004488	Homo sapiens glycoprotein V (platelet) (GP5), mRNA
NM_004485	Homo sapiens guanine nucleotide binding protein 4 (GNG4), mRNA
NM_004122	Homo sapiens growth hormone secretagogue receptor (GHSR), mRNA
NM_004479	Homo sapiens fucosyltransferase 7 (alpha (1,3) fucosyltransferase) (FUT7), mRNA
NM_004438	Homo sapiens EphA4 (EPHA4), mRNA
NM_004094	Homo sapiens eukaryotic translation initiation factor 2, subunit 1 (alpha, 35kD) (EIF2S1), mRNA
NM_004681	Homo sapiens eukaryotic translation initiation factor 1A, Y chromosome (EIF1AY), mRNA
NM_004226	Homo sapiens serine/threonine kinase 17b (apoptosis-inducing) (STK17B), mRNA
NM_004792	Homo sapiens peptidyl-prolyl isomerase G (cyclophilin G) (PPIG), mRNA
NM_004831	Homo sapiens cofactor required for Sp1 transcriptional activation, subunit 7 (70kD) (CRSP7), mRNA
NM_004269	Homo sapiens cofactor required for Sp1 transcriptional activation, subunit 8 (34kD) (CRSP8), mRNA
NM_004270	Homo sapiens cofactor required for Sp1 transcriptional activation, subunit 9 (33kD) (CRSP9), mRNA
NM_004232	Homo sapiens STAT induced STAT inhibitor-4 (CIS4), mRNA
NM_004882	Homo sapiens CBF1 interacting corepressor (CIR), mRNA
NM_004198	Homo sapiens cholinergic receptor, nicotinic, alpha polypeptide 6 (CHRNA6), mRNA
NM_004825	Homo sapiens chromodomain protein, Y chromosome, 2 (CDY2), mRNA
NM_004351	Homo sapiens Cas-Br-M (murine) ectropic retroviral transforming sequence b (CBLB), mRNA
NM_004054	Homo sapiens complement component 3a receptor 1 (C3AR1), mRNA
	1,2

NM_004899	Homo sapiens brain and reproductive organ-expressed (TNFRSF1A modulator) (BRE), mRNA
NM_004889	Homo sapiens ATP synthase, H+ transporting, mitochondrial F0 complex, subunit f, isoform 2 (ATP5J2), mRNA
NM 004890	Homo sapiens sperm associated antigen 7 (SPAG7), mRNA
NM 004908	Homo sapiens pre-T/NK cell associated protein (6H9A), mRNA
NM 003406	Homo sapiens tyrosine 3-monooxygenase/tryptophan 5-monooxygenase
_	activation protein, zeta polypeptide (YWHAZ), mRNA
NM_003574	Homo sapiens VAMP (vesicle-associated membrane protein)-associated protein A (33kD) (VAPA), mRNA
NM_001073	Homo sapiens UDP glycosyltransferase 2 family, polypeptide B11 (UGT2B11), mRNA
NM_003300	Homo sapiens TNF receptor-associated factor 3 (TRAF3), mRNA
NM_003297	Homo sapiens nuclear receptor subfamily 2, group C, member 1 (NR2C1), mRNA
NM 003212	Homo sapiens teratocarcinoma-derived growth factor 1 (TDGF1), mRNA
NM 003763	Homo sapiens syntaxin 16 (STX16), mRNA
NM 003955	Homo sapiens STAT induced STAT inhibitor 3 (SSI-3), mRNA
NM_003693	Homo sapiens acetyl LDL receptor; SREC=scavenger receptor expressed by endothelial cells (SREC), mRNA
NM 003563	Homo sapiens speckle-type POZ protein (SPOP), mRNA
NM 003578	Homo sapiens sterol O-acyltransferase 2 (SOAT2), mRNA
NM 003099	Homo sapiens sorting nexin 1 (SNX1), mRNA
NM 003095	Homo sapiens small nuclear ribonucleoprotein polypeptide F (SNRPF), mRNA
NM 003091	Homo sapiens small nuclear ribonucleoprotein polypeptides B and B1 (SNRPB),
	mRNA
NM_003086	Homo sapiens small nuclear RNA activating complex, polypeptide 4, 190kD (SNAPC4), mRNA
NM_003084	Homo sapiens small nuclear RNA activating complex, polypeptide 3, 50kD (SNAPC3), mRNA
NM_003825	Homo sapiens synaptosomal-associated protein, 23kD (SNAP23), mRNA
NM_003983	Homo sapiens solute carrier family 7 (cationic amino acid transporter, y+ system), member 6 (SLC7A6), mRNA
NM_003916	Homo sapiens adaptor-related protein complex 1, sigma 2 subunit (AP1S2), mRNA
NM_003896	Homo sapiens sialyltransferase 9 (CMP-NeuAc:lactosylceramide alpha-2,3-sialyltransferase; GM3 synthase) (SIAT9), mRNA
NM 003769	Homo sapiens splicing factor, arginine/serine-rich 9 (SFRS9), mRNA
NM 003016	Homo sapiens splicing factor, arginine/serine-rich 2 (SFRS2), mRNA
NM_003161	Homo sapiens ribosomal protein S6 kinase, 70kD, polypeptide 1 (RPS6KB1), mRNA
NM_003708	Homo sapiens microsomal NAD+-dependent retinol dehydrogenase 4 (RODH-4), mRNA
NM 002933	Homo sapiens ribonuclease, RNase A family, 1 (pancreatic) (RNASE1), mRNA
NM_002919	Homo sapiens regulatory factor X, 3 (influences HLA class II expression) (RFX3), mRNA
NM 002865	Homo sapiens RAB2, member RAS oncogene family (RAB2), mRNA
NM 002849	Homo sapiens protein tyrosine phosphatase, receptor type, R (PTPRR), mRNA
NM 002822	Homo sapiens protein tyrosine kinase 9 (PTK9), mRNA
NM_002812	Homo sapiens proteasome (prosome, macropain) 26S subunit, non-ATPase, 8 (PSMD8), mRNA
NM_002808	Homo sapiens proteasome (prosome, macropain) 26S subunit, non-ATPase, 2

	(PSMD2), mRNA
NM_002816	Homo sapiens proteasome (prosome, macropain) 26S subunit, non-ATPase, 12
	(1 SNID 12), IIIXIVA
NM_002814	1 The substitute of the state o
	_   (I BIVID IU), IIIKIVA
NM_002789	The state of the s
	(I BIVIA4), IIIKIVA
NM_002787	Homo sapiens proteasome (prosome, macropain) subunit, alpha type, 2
NM_000951	Homo sapiens proline-rich Gla (G-carboxyglutamic acid) polypeptide 2
377	
NM_000950	Homo sapiens proline-rich Gla (G-carboxyglutamic acid) polypeptide 1
375	(1001), IIINA
NM_002750	Homo sapiens mitogen-activated protein kinase 8 (MAPK8), mRNA
NM_003981	Tionio sapiens protein regulator of cytokinesis 1 (DDC1) DATA
NM_002717	1101110 Sapiens protein phosphatase 2 (formerly 2A) regulaters 1 in D. (20)
NM_002707	Homo sapiens protein phosphatase 1G (formerly 2C), magnetism 1
	Barrante isololiii (1 1 MICI). IIIR NA
NM_003620	Homo sapiens protein phosphatase 1D magnesium-dependent, doltain 5
37.	
NM_003625	Homo sapiens protein tyrosine phosphatase, receptor type, f polypeptide
370.6	.   (* ** ** /) IIIOI (IIII   DIDIEII   HINTIN   Alpha // (DDET & 7) pat A
NM 002698	Tiomo sapiens POU domain, class 2, transcription factor 2 (POUDE2) - PALA
NM_002687	
NM_003662	Tiono sapiens Pinn (PIR), mRNA
NM_002647	Homo sapiens phosphoinositide-3-kinase class 3 (PIX 3C2) PNA
NM_000286	Tiomo sapiens peroxisomal biogenesis factor 12 (DEV12) DATA
NM_002861	Tiomo sapiens phosphate cytidylyltransferase 2 ethanologia (DCVVI)
NM_002567	1 220 MO Suproits prostatic difficition protein (PRP) and XI A
NM_003899	Homo sapiens Rho guanine nucleotide exchange factor (GEE) 7 (A DIJGERT)
ND 6 0007 60	
NM_002563	Homo sapiens purinergic receptor P2Y, G-protein coupled, 1 (P2RY1), mRNA
NM_000913	1 - 1 - 1 - 1 - 1 - 1 - 1 - 1 - 1 - 1 -
NM_002493	nomo sapiens NADH dehydrogenase (ubiquinone) 1 beta gubogenalari 6 (171 p.
ND ( 000400	1 DOI DOI, MILLINA
NM_002492	Homo sapiens NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 5 (16kD, SGDH) (NDUERS) mPNA
ND 4 002400	TODAY (INDOING), IIICINA
NM_002489	Homo sapiens NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 4 (9kD, MLRO) (NDIFA4), mPNIA
NM_003684	
11111_003084	Homo sapiens MAP kinase-interacting serine/threonine kinase 1 (MKNK1),
NM_003784	med 41.
INIM_003784	Homo sapiens serine (or cysteine) proteinase inhibitor, clade B (ovalbumin),
NTM 000000	Member / (BERTIND/), MRNA
NM_002333	Homo sapiens low density lipoprotein receptor-related protein 3 (LRP3), mRNA
NM_002285	
VM_002213	220 me Sapiciis integrin, deta 5 (11(4R5) me Nia
VM_003971	nomo sapiens sperm associated antigen 9 (SDA GO) - DNA
VM_002157	Tiomo sapiens neat shock 10kD protein 1 (chaperonin 10) (USDE1)
VM_001521	suprems general transcription factor IIIC nolymentide 2 Chata and
VM_001516	
ATAT_001210	Homo sapiens general transcription factor IIH, polypeptide 3 (34kD subunit)
	F-5F-Passo (STAD SHOUMIL)

	(GTF2H3), mRNA
NM_003910	Homo sapiens maternal G10 transcript (G10), mRNA
NM_001969	Homo sapiens eukaryotic translation initiation factor 5 (EIF5), mRNA
NM_003751	Homo sapiens eukaryotic translation initiation factor 3, subunit 9 (eta, 116kD) (EIF3S9), mRNA
NM_003755	Homo sapiens eukaryotic translation initiation factor 3, subunit 4 (delta, 44kD) (EIF3S4), mRNA
NM_003756	Homo sapiens eukaryotic translation initiation factor 3, subunit 3 (gamma, 40kD) (EIF3S3), mRNA
NM_001414	Homo sapiens eukaryotic translation initiation factor 2B, subunit 1 (alpha, 26kD) (EIF2B1), mRNA
NM_001412	Homo sapiens eukaryotic translation initiation factor 1A (EIF1A), mRNA
NM 003566	Homo sapiens early endosome antigen 1, 162kD (EEA1), mRNA
NM 001957	Homo sapiens endothelin receptor type A (EDNRA), mRNA
NM 001936	Homo sapiens dipeptidylpeptidase VI (DPP6), mRNA
NM 003648	Homo sapiens diacylglycerol kinase, delta (130kD) (DGKD), mRNA
NM 001921	Homo sapiens dCMP deaminase (DCTD), mRNA
NM_003590	Homo sapiens cullin 3 (CUL3), mRNA
NM 003592	Homo sapiens cullin 1 (CUL1), mRNA
NM 001207	Homo sapiens basic transcription factor 3 (BTF3), mRNA
NM 001191	Homo sapiens BCL2-like 1 (BCL2L1), mRNA
NM 001689	Homo sapiens ATP synthase, H+ transporting, mitochondrial F0 complex,
14141_001085	subunit c (subunit 9) isoform 3 (ATP5G3), mRNA
NM_001688	Homo sapiens ATP synthase, H+ transporting, mitochondrial F0 complex,
14141_001000	subunit b, isoform 1 (ATP5F1), mRNA
NM_003664	Homo sapiens adaptor-related protein complex 3, beta 1 subunit (AP3B1),
14141_003004	mRNA
NM 058168	Homo sapiens gene differentially expressed in prostate (GDEP), mRNA
NM 058222	Homo sapiens tectorin beta (TECTB), mRNA
NM 058192	Homo sapiens ribosomal large subunit pseudouridine synthase C like (RLUCL),
	mRNA
NM_058190	Homo sapiens chromosome 21 open reading frame 70 (C21orf70), mRNA
NM_058189	Homo sapiens chromosome 21 open reading frame 69 (C21orf69), mRNA
NM_058186	Homo sapiens chromosome 21 open reading frame 11 (C21orf11), mRNA
NM_058184	Homo sapiens chromosome 21 open reading frame 42 (C21orf42), mRNA
NM_058182	Homo sapiens chromosome 21 open reading frame 51 (C21orf51), mRNA
NM_058180	Homo sapiens chromosome 21 open reading frame 58 (C21orf58), mRNA
NM_058173	Homo sapiens small breast epithelial mucin (LOC118430), mRNA
NM_058172	Homo sapiens capillary morphogenesis protein 2 (CMG2), mRNA
NM_017884	Homo sapiens PIN2-interacting protein 1 (PINX1), mRNA
NM_054021	Homo sapiens G protein-coupled receptor 101 (GPR101), mRNA
NM_053280	Homo sapiens h-Shippo 1 (LOC113746), mRNA
NM_003449	Homo sapiens tripartite motif-containing 26 (TRIM26), mRNA
NM_052939	Homo sapiens Fc receptor-like protein 3 (FCRH3), mRNA
NM_052938	Homo sapiens Fc receptor-like protein 1 (FCRH1), mRNA
NM_052872	Homo sapiens interleukin 17F (IL17F), mRNA
NM_024011	Homo sapiens cell division cycle 2-like 2 (CDC2L2), transcript variant 1, mRNA
NM_033621	Homo sapiens cell division cycle 2-like 2 (CDC2L2), transcript variant 10,
	mRNA
NM 033537	Homo sapiens cell division cycle 2-like 2 (CDC2L2), transcript variant 9, mRNA
NM 033536	Homo sapiens cell division cycle 2-like 2 (CDC2L2), transcript variant 8, mRNA
NM 033534	Homo sapiens cell division cycle 2-like 2 (CDC2L2), transcript variant 7, mRNA
1111 000004	220

	2 (CD COLO) to resint verient 6 mRNA
NM 033532	Homo sapiens cell division cycle 2-like 2 (CDC2L2), transcript variant 6, mRNA
NM 033531	Homo sapiens cell division cycle 2-like 2 (CDC2L2), transcript variant 5, mRNA  Homo sapiens cell division cycle 2-like 2 (CDC2L2), transcript variant 4, mRNA
NM 033529	Homo sapiens cell division cycle 2-like 2 (CDC2L2), transcript variant 4, mRNA  Homo sapiens cell division cycle 2-like 2 (CDC2L2), transcript variant 4, mRNA
NM 033528	Homo sapiens cell division cycle 2-like 2 (CDC2L2), transcript variant 3, mRNA  Homo sapiens cell division cycle 2-like 2 (CDC2L2), transcript variant 3, mRNA
NM 033527	Homo sapiens cell division cycle 2-like 2 (CDC2L2), transcript variant 2, micror
NM 006629	Homo seniens zinc finger protein 271 (ZNF2/1), mRNA
NM 015294	Homo sapiens tripartite motif-containing 37 (TRIM37), mRNA
NM 033132	Homo capiens zinc family member 5 protein (ZICS), mRNA
NM 033108	Home saniens heat shock transcription factor 2-like (LOC80014), IIIKNA
NM 033106	Homo saniens galanin-like peptide precursor (LOC85569), mRNA
NM 033105	Homo sapiens beta cysteine string protein (LOC85479), mRNA
NM 033104	Homo saniens stonin 2 (LOC85439), mRNA
NM 033102	Thems comions prostein protein (LOC85414), mRNA
NM 003823	Homo saniens tumor necrosis factor receptor superfamily, member 60, decoy
14141_005025	(TNFR SF6B), transcript variant M68E, mRNA
NM 006470	Homo sapiens tripartite motif-containing 16 (TRIM16), mRNA
NM 032606	Homo sapiens calcyphosine (LOC84698), mRNA
NM 032595	Homo saniens neurabin II (LOC84687), mRNA
NM 032584	Hame senions zinc finger protein 347 (ZNF347), mRNA
NM_032576	Homo sapiens lipopolysaccaride-specific response 5-like protein (LOC84663),
14141_052570	mDNA
NM_032518	Homo sapiens collagen-like Alzheimer amyloid plaque component precursor
1411_052510	(I OC84570), mRNA
NM 032509	Homo sapiens RNA binding protein (LOC84549), mRNA
NM 032484	Homo seriens hypothetical protein (LOC84514), mKNA
NM 032389	Homo sapiens zinc finger protein 289, ID1 regulated (ZNF289), mRNA
NM 031918	Homo sapiens Kruppel-like factor 16 (KLF16), mRNA
NM 031463	Homo sapiens steroid dehydrogenase-like (LOC83693), mRNA
NM 031461	III and serious Cocoo Crisp (LOC83690), mRNA
NM_031417	Homo sapiens Cocoacrisp (Locososo), mention sapiens MAP/microtubule affinity-regulating kinase like 1 (MARKL1),
112.2_00	mRNA
NM 030791	Homo sapiens sphingosine-1-phosphatase (LOC81537), mRNA
NM_024670	Homo sapiens suppressor of variegation 3-9 (Drosophila) homolog 2;
	hymothetical protein FLI23414 (SUV39H2), mRNA
NM_003414	Homo sapiens zinc finger protein 267 (ZNF267), transcript variant 498723,
	DNIA
NM_023945	Homo sapiens membrane-spanning 4-domains, subfamily A, member 5
	(MCAAS) mDNA
NM_023014	Homo sapiens hypothetical protein similar to preferentially expressed antigen of
_	1
NM_023013	Homo sapiens hypothetical protein similar to preferentially expressed antigen of
	melanoma (I OC65121), mRNA
NM_022357	Homo sapiens putative metallopeptidase (family M19) (LOC64180), mRNA
NM_022355	Home saniers putative dipeptidase (LOC64174), mRNA
NM_022353	Homo sapiens putative sialoglycoprotease type 2 (LOC64172), mRNA
NM 022345	Homo sapiens uterine-derived 14 kDa protein (LOC64150), mRNA
NM 022343	Homo sapiens 17kD fetal brain protein (LOC64148), mRNA
NM 022340	Homo sapiens FYVE-finger-containing Rab5 effector protein rabenosyn-5
_	(I OC64145) mRNA
NM_021932	C PUDODA A CE 1097170 // OCO00/01
	mRNA Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 35 (DDX35),
NM 021931	TEADIL (Am Gly Ale Am/Hig) how notworning to (DDA)

[	mRNA
NM 021632	Homo sapiens zinc-finger protein ZBRK1 (ZBRK1), mRNA
NM 021630	Homo sapiens PDZ-LIM protein mystique (LOC59346), mRNA
NM 019591	Homo sapiens zinc finger protein 26 (KOX 20) (ZNF26), mRNA
NM 018675	Homo sapiens zinc finger protein 302 (ZNF302), mRNA
NM 021226	Homo sapiens hypothetical protein from clones 23549 and 23762 (LOC58504),
1111_021220	mRNA
NM_021211	Homo sapiens transposon-derived Buster1 transposase-like protein (LOC58486),
	mRNA
NM 021186	Homo sapiens zona pellucida glycoprotein 4 (ZP4), mRNA
NM 020903	Homo sapiens ubiquitin-specific processing protease (LOC57663), mRNA
NM 020666	Homo sapiens CDC-like kinase 4 (CLK4), mRNA
NM 020421	Homo sapiens hypothetical protein (LOC57143), mRNA
NM 020140	Homo sapiens putative 47 kDa protein (LOC56899), mRNA
NM 016305	Homo sapiens synovial sarcoma translocation gene on chromosome 18-like 2
_	(SS18L2), mRNA
NM_016417	Homo sapiens clone FLB4739 (LOC51218), mRNA
NM 020467	Homo sapiens hypothetical protein from clone 643 (LOC57228), mRNA
NM_020389	Homo sapiens putative capacitative calcium channel (trp7), mRNA
NM 020385	Homo sapiens XPMC2 protein (LOC57109), mRNA
NM 020381	Homo sapiens candidate tumor suppressor protein (LOC57107), mRNA
NM 020372	Homo sapiens organic cation transporter (LOC57100), mRNA
NM 020158	Homo sapiens exosome component Rrp46 (RRP46), mRNA
NM 020147	Homo sapiens hypothetical protein from EUROIMAGE 511235 (LOC56906),
L	mRNA
NM_020154	Homo sapiens chromosome 11 hypothetical protein ORF3 (LOC56851), mRNA
NM_019613	Homo sapiens hypothetical protein 628 (LOC56270), mRNA
NM_019059	Homo sapiens 6.2 kd protein (LOC54543), mRNA
NM_019037	Homo sapiens exosome component Rrp41 (FLJ20591), mRNA
NM_018579	Homo sapiens mitochondrial solute carrier (LOC51312), mRNA
NM_018485	Homo sapiens G protein-coupled receptor C5L2 (LOC55868), mRNA
NM_018479	Homo sapiens uncharacterized hypothalamus protein HCDASE (LOC55862),
	mRNA
NM_018447	Homo sapiens 30 kDa protein (LOC55831), mRNA
NM_018443	Homo sapiens zinc finger protein 302 (ZNF302), mRNA
NM_018430	Homo sapiens hypothetical protein (LOC55815), mRNA
NM_018402	Homo sapiens interleukin 26 (IL26), mRNA
NM_017692	Homo sapiens aprataxin (APTX), mRNA
NM_018171	Homo sapiens hypothetical protein FLJ10659 (FLJ10659), mRNA
NM_017530	Homo sapiens hypothetical protein LOC55565 (LOC55565), mRNA
NM_013385	Homo sapiens pleckstrin homology, Sec7 and coiled/coil domains 4 (PSCD4), mRNA
NM_016651	Homo sapiens heptacellular carcinoma novel gene-3 protein (LOC51339), mRNA
NM 016955	Homo sapiens soluble liver antigen/liver pancreas antigen (LOC51091), mRNA
NM 016422	Homo sapiens C3HC4-like zinc finger protein (ZFP26), mRNA
NM_016520	Homo sapiens hepatocellular carcinoma-associated antigen 59 (LOC51759), mRNA
NM_016275	Homo sapiens selenoprotein T (LOC51714), mRNA
NM 016242	Homo sapiens endomucin-2 (LOC51705), mRNA
NM 016233	Homo sapiens peptidylarginine deiminase type III (LOC51702), mRNA
NM 016209	Homo sapiens unknown (LOC51693), mRNA
1111_010207	1 azeme especie diliciowii (LOCS1000), ilitari

	7 0 C 5 1 (72) PN A
VM 016140	Homo sapiens brain specific protein (LOC51673), mRNA
Th ( 016107	Homo saniens zinc finger RNA binding protein (ZFK), IIIKNA
NTA 016009	Homo saniens HSPC040 protein (LOC51660), mRNA
	TICDC027 protein (LOC51659), MKNA
NM_016086	Homo sapiens HSPC037 protein (LSCSTeSS), Homo sapiens map kinase phosphatase-like protein MK-STYX (LOC51657),
_	mR N A
NM 016061	Homo sapiens CGI-127 protein (LOC51646), mRNA
NM 016039	Homo sapiens CGI-99 protein (LOC51637), mRNA
NM 016029	Homo saniens CGI-86 protein (LOC51635), mRNA
NM 016024	Homo saniens CGI-79 protein (LOC51634), mRNA
NM 016019	Homo saniens CGI-74 protein (LOC51631), mRNA
NM 015964	Homo sapiens brain specific protein (LOC51673), mRNA
NM 015939	CCT 00 meetsin (1 () ('\$1603) MK NA
NM 016647	rx see maganahymal stem cell protein DSCD/3 (LOC31337), indivi
NM 016646	Homo sapiens mesenchymal stem cell protein DSCD28 (LOCS1330), mid vi
NM 016632	Homo saniens ARF protein (LOC51326), mRNA
NM 016629	Homo saniens hypothetical protein (LOC51323), mRNA
NM 016627	Homo caniens hypothetical protein (LOC51321), mkNA
NM 016626	Homo saniens hypothetical protein (LOCS1320), mkNA
	Homo sapiens hypothetical protein (LOC51315), mRNA
NM 016618	Homo sapiens NM23-H8 (LOC51314), mRNA
NM_016616	Trans senions AD021 protein (LOC51313), MKNA
NM_016613	Home conjens mitochondrial solute carrier (LOC51312), Illinia
NM_016612	Homo sapiens FK506 binding protein precursor (LOC51303), mRNA
NM_016594	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
NM 016562	TI are agricus complement ("Ir-like profemase precuisor, (LOC51279), me uz
NM_016546	Homo sapiens apoptosis-related protein PNAS-1 (LOC51275), mRNA
NM_016534	Lucro saniers F2F-like protein (LOC51270), mRNA
NM_016521	Home conjens C-type lectin-like receptor-1 (LOC51207), Illicina
NM_016511	Homo sapiens C-type lectin-like receptor-2 (LOC51266), mRNA  Homo sapiens C-type lectin-like receptor-2 (LOC51266), mRNA
NM_016509	Homo sapiens hypothetical protein (LOC31237), mkNA
NM_016496	Homo sapiens hypothetical protein (LOC51255), mRNA  Homo sapiens hypothetical protein (LOC51255), mRNA
NM_016494	Homo sapiens hypothetical protein (LOC51248), mRNA  Homo sapiens hypothetical protein (LOC51248), mRNA
NM_016484	Homo sapiens hypothetical protein (LOC51242), mRNA  Homo sapiens hypothetical protein (LOC51242), mRNA
NM 016471	Homo sapiens hypothetical protein (LOC51240), mRNA  Homo sapiens hypothetical protein (LOC51240), mRNA
NM 016467	TYiong hymothetical protein (1.000) 234). HINNA
NM_016454	Homo sapiens COPZ2 for nonclathrin coat protein zeta-COP (LOC51226),
NM_016429	
	mRNA Homo sapiens HOM-TES-85 tumor antigen (LOC51213), mRNA
NM_016383	Homo sapiens diferentiation-related protein dif13 (LOC51212), mRNA
NM_016380	The state of the substance (10) 11 (11) MKINA
NM_016364	T 1
NM_016339	T 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
NM_016338	Homo sapiens zinc finger protein ANC 2H01 (LOC51193), mRNA  Homo sapiens zinc finger protein ANC 2H01 (LOC51193), mRNA
NM_016331	
NM_016311	- I - b conbodiecter 2IDD2-IV-
NM_016256	Homo sapiens N-acetylgiucosainine-1-phosphodiosis arpan a
	acetylglucosaminidase (LOC51172), mRNA  Homo sapiens protein kinase C and casein kinase substrate in neurons 3
NM_016223	Homo sapiens protein kinase C and casem kinase substate at the most of the case of the cas
	(PACSIN3), mRNA
NM_016202	Homo sapiens LDL induced EC protein (LOC51157), mRNA  Homo sapiens LDL induced EC protein (LOC51149), mRNA
NM_016175	
NM 016162	Homo sapiens candidate tumor suppressor p33 ING1 homolog (LOC51147),

	DNIA
ND4 016150	mRNA
NM_016158	Homo sapiens erythrocyte transmembrane protein (LOC51145), mRNA
NM_016142	Homo sapiens steroid dehydrogenase homolog (LOC51144), mRNA
NM_016141	Homo sapiens dynein light chain-A (LOC51143), mRNA
NM_016125	Homo sapiens PTD016 protein (LOC51136), mRNA
NM_016121	Homo sapiens NY-REN-45 antigen (LOC51133), mRNA
NM_016102	Homo sapiens tripartite motif-containing 17 (TRIM17), mRNA
NM_016038	Homo sapiens CGI-97 protein (LOC51119), mRNA
NM_016035	Homo sapiens CGI-92 protein (LOC51117), mRNA
NM_016026	Homo sapiens CGI-82 protein (LOC51109), mRNA
NM_016010	Homo sapiens CGI-62 protein (LOC51101), mRNA
NM_016001	Homo sapiens CGI-48 protein (LOC51096), mRNA
NM_015996	Homo sapiens CGI-40 protein (LOC51092), mRNA
NM_015978	Homo sapiens putative protein-tyrosine kinase (LOC51086), mRNA
NM_015962	Homo sapiens CGI-35 protein (LOC51077), mRNA
NM_015960	Homo sapiens CGI-32 protein (LOC51076), mRNA
NM_015957	Homo sapiens CGI-29 protein (LOC51074), mRNA
NM_015954	Homo sapiens CGI-26 protein (LOC51071), mRNA
NM_015917	Homo sapiens glutathione S-transferase subunit 13 homolog (LOC51064), mRNA
NM 015913	Homo sapiens hypothetical protein (LOC51060), mRNA
NM 015912	Homo sapiens hypothetical protein (LOC51059), mRNA
NM 015911	Homo sapiens hypothetical protein (LOC51058), mRNA
NM 015907	Homo sapiens leucine aminopeptidase (LOC51056), mRNA
NM 015883	Homo sapiens clone 1900 unknown protein (LOC51049), mRNA
NM 015872	Homo sapiens kruppel-related zinc finger protein hcKrox (LOC51043), mRNA
NM 015871	Homo sapiens zinc finger protein (LOC51042), mRNA
NM 016072	Homo sapiens CGI-141 protein (LOC51026), mRNA
NM 016068	Homo sapiens CGI-135 protein (LOC51024), mRNA
NM 016053	Homo sapiens CGI-116 protein (LOC51019), mRNA
NM 016046	Homo sapiens homolog of yeast exosomal core protein CSL4 (CSL4), mRNA
NM 016042	Homo sapiens exosome component Rrp40 (RRP40), mRNA
NM 015944	Homo sapiens CGI-14 protein (LOC51005), mRNA
NM 016060	Homo sapiens CGI-125 protein (LOC51003), mRNA
NM_016482	Homo sapiens hepatocellular carcinoma-associated antigen 59 (LOC51759), mRNA
NM_014681	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 34 (DDX34),
	mRNA
NM_014415	Homo sapiens zinc finger protein (ZNF-U69274), mRNA
NM_014579	Homo sapiens zinc transporter (ZIP2), mRNA
NM_014347	Homo sapiens zinc finger protein (ZF5128), mRNA
NM_007146	Homo sapiens zinc finger protein 161 (ZNF161), mRNA
NM_006626	Homo sapiens zinc finger protein with interaction domain (ZID), mRNA
NM_006336	Homo sapiens ZYG homolog (ZYG), mRNA
NM_006138	Homo sapiens membrane-spanning 4-domains, subfamily A, member 3 (hematopoietic cell-specific) (MS4A3), mRNA
NM 005741	Homo sapiens zinc finger protein 263 (ZNF263), mRNA
NM_000227	Homo sapiens laminin, alpha 3 (nicein (150kD), kalinin (165kD), BM600
	(150kD), epilegrin) (LAMA3), mRNA
NM_000423	Homo sapiens keratin 2A (epidermal ichthyosis bullosa of Siemens) (KRT2A), mRNA
NM_000659	Homo sapiens autoimmune regulator (automimmune polyendocrinopathy

	candidiasis ectodermal dystrophy) (AIRE), transcript variant 3, mRNA
NM_000658	Homo sapiens autoimmune regulator (automimmune polyendocrinopathy
	candidiasis ectodermal dystrophy) (AIRE), transcript variant AIRE-2, mRNA
NM_000383	Homo sapiens autoimmune regulator (automimmune polyendocrinopathy
	candidiasis ectodermal dystrophy) (AIRE), transcript variant AIRE-1, mRNA
NM_003451	Homo sapiens zinc finger protein 177 (ZNF177), mRNA
NM_003419	Homo sapiens zinc finger protein 345 (ZNF345), mRNA
NM_003407	Homo sapiens zinc finger protein 36, C3H type, homolog (mouse) (ZFP36),
	mRNA
NM_001519	Homo sapiens BRF1 homolog, subunit of RNA polymerase III transcription
	initiation factor IIIB (S.cerevisiae) (BRF1), mRNA
NM_000157	Homo sapiens glucosidase, beta; acid (includes glucosylceramidase) (GBA),
	mRNA
NM_057178	Homo sapiens fring (LOC117584), mRNA
NM_057177	Homo sapiens amyotrophic lateral sclerosis 2 (juvenile) chromosome region,
_	candidate 19 (ALS2CR19), mRNA
NM_058178	Homo sapiens neuronal pentraxin receptor (NPTXR), transcript variant 2, mRNA
NM 014293	Homo sapiens neuronal pentraxin receptor (NPTXR), transcript variant 1, mRNA
NM 012223	Homo sapiens myosin IB (MYO1B), mRNA
NM 015277	Homo sapiens neural precursor cell expressed, developmentally down-regulated
_	4-like (NEDD4L), mRNA
NM 015074	Homo sapiens kinesin family member 1B (KIF1B), mRNA
NM 032591	Homo sapiens solute carrier family 9 (sodium/hydrogen exchanger), isoform 7
_	(SLC9A7), mRNA
NM 014208	Homo sapiens dentin sialophosphoprotein (DSPP), mRNA
NM 014693	Homo sapiens endothelin converting enzyme 2 (ECE2), mRNA
NM 005461	Homo sapiens v-maf musculoaponeurotic fibrosarcoma oncogene homolog B
_	(avian) (MAFB), mRNA
NM_030761	Homo sapiens wingless-type MMTV integration site family, member 4 (WNT4),
	mRNA
NM_032642	Homo sapiens wingless-type MMTV integration site family, member 5B
	(WNT5B), transcript variant 1, mRNA
NM_030775	Homo sapiens wingless-type MMTV integration site family, member 5B
	(WNT5B), transcript variant 2, mRNA
NM_003392	Homo sapiens wingless-type MMTV integration site family, member 5A
	(WNT5A), mRNA
NM_057168	Homo sapiens wingless-type MMTV integration site family, member 16
	(WNT16), transcript variant 1, mRNA
NM_016087	Homo sapiens wingless-type MMTV integration site family, member 16
	(WNT16), transcript variant 2, mRNA
NM_012101	Homo sapiens tripartite motif-containing 29 (TRIM29), transcript variant 1,
	mRNA
NM_058193	Homo sapiens tripartite motif-containing 29 (TRIM29), transcript variant 2,
	mRNA
NM_000983	Homo sapiens ribosomal protein L22 (RPL22), mRNA
NM_058248	Homo sapiens DNase II-like acid DNase (DLAD), transcript variant 2, mRNA
NM_021233	Homo sapiens DNase II-like acid DNase (DLAD), transcript variant 1, mRNA
NM_058175	Homo sapiens collagen, type VI, alpha 2 (COL6A2), transcript variant 2C2a',
	mRNA
NM_058174	Homo sapiens collagen, type VI, alpha 2 (COL6A2), transcript variant 2C2a,
	mRNA
NM_001849	Homo sapiens collagen, type VI, alpha 2 (COL6A2), transcript variant 2C2,

	mRNA
NM 003312	Homo sapiens thiosulfate sulfurtransferase (rhodanese) (TST), mRNA
	<del>  </del>
NM_020731	Homo sapiens dioxin receptor repressor (AHRR), mRNA
NM_053049	Homo sapiens stresscopin (SPC), mRNA
NM_052834	Homo sapiens WD repeat domain 7 (WDR7), transcript variant 2, mRNA
NM_015285	Homo sapiens WD repeat domain 7 (WDR7), transcript variant 1, mRNA
NM_000507	Homo sapiens fructose-1,6-bisphosphatase 1 (FBP1), mRNA
NM_002581	Homo sapiens pregnancy-associated plasma protein A (PAPPA), mRNA
NM_000968	Homo sapiens ribosomal protein L4 (RPL4), mRNA
NM_005061	Homo sapiens ribosomal protein L3-like (RPL3L), mRNA
NM_030811	Homo sapiens mitochondrial ribosomal protein S26 (MRPS26), nuclear gene encoding mitochondrial protein, mRNA
NM 022497	Homo sapiens mitochondrial ribosomal protein S25 (MRPS25), nuclear gene
-	encoding mitochondrial protein, mRNA
NM_053023	Homo sapiens zinc finger protein homologous to Zfp91 in mouse (ZFP91), mRNA
NM 052826	Homo sapiens WD repeat domain 6 (WDR6), transcript variant 2, mRNA
NM 052825	Homo sapiens WD repeat domain 6 (WDR6), transcript variant 2, mRNA
NM 052821	Homo sapiens WD repeat domain 5 (WDR5), transcript variant 2, mRNA
NM 017588	Homo sapiens WD repeat domain 5 (WDR5), transcript variant 1, mRNA
NM 052990	Homo sapiens WD repeat domain 10 (WDR10), transcript variant 4, mRNA
NM 052989	Homo sapiens WD repeat domain 10 (WDR10), transcript variant 4, mRNA
NM 052985	Homo sapiens WD repeat domain 10 (WDR10), transcript variant 2, mRNA
NM 018262	Homo sapiens WD repeat domain 10 (WDR10), transcript variant 1, mRNA
NM_031902	Homo sapiens mitochondrial ribosomal protein S5 (MRPS5), nuclear gene
14141_051702	encoding mitochondrial protein, mRNA
NM 015969	Homo sapiens mitochondrial ribosomal protein S17 (MRPS17), nuclear gene
14112_010505	encoding mitochondrial protein, mRNA
NM_016065	Homo sapiens mitochondrial ribosomal protein S16 (MRPS16), nuclear gene
_	encoding mitochondrial protein, mRNA
NM_031280	Homo sapiens mitochondrial ribosomal protein S15 (MRPS15), nuclear gene
_	encoding mitochondrial protein, mRNA
NM 022839	Homo sapiens mitochondrial ribosomal protein S11 (MRPS11), nuclear gene
_	encoding mitochondrial protein, mRNA
NM 016034	Homo sapiens mitochondrial ribosomal protein S2 (MRPS2), nuclear gene
	encoding mitochondrial protein, mRNA
NM_016070	Homo sapiens mitochondrial ribosomal protein S23 (MRPS23), nuclear gene
	encoding mitochondrial protein, mRNA
NM_020191	Homo sapiens mitochondrial ribosomal protein S22 (MRPS22), nuclear gene
	encoding mitochondrial protein, mRNA
NM_018135	Homo sapiens mitochondrial ribosomal protein S18A (MRPS18A), nuclear gene
	encoding mitochondrial protein, mRNA
NM_021996	Homo sapiens Forssman glycolipid synthetase (FS), mRNA
NM_052815	Homo sapiens immediate early response 3 (IER3), transcript variant long,
L	mRNA
NM_003897	Homo sapiens immediate early response 3 (IER3), transcript variant short, mRNA
NM 053013	Homo sapiens enolase 3, (beta, muscle) (ENO3), transcript variant 2, mRNA
NM 001976	Homo sapiens enolase 3, (beta, muscle) (ENO3), transcript variant 1, mRNA
NM 048368	Homo sapiens CTD (carboxy-terminal domain, RNA polymerase II, polypeptide
	A) phosphatase, subunit 1 (CTDP1), transcript variant FCP1b, mRNA
NM_004715	Homo sapiens CTD (carboxy-terminal domain, RNA polymerase II, polypeptide

	A) phosphatase, subunit 1 (CTDP1), transcript variant FCP1a, mRNA
	Homo sapiens collagen, type V, alpha 3 (COL5A3), mRNA
NM_015719	Homo sapiens collagen, type V, alpha 2 (COL5A2), mRNA  Homo sapiens collagen, type V, alpha 2 (COL5A2), mRNA
NM_000393	Homo sapiens collagen, type V, alpha 1 (COL5A1), mRNA
NM_000093	Homo sapiens collagen, type V, appla 1 (COESTE), MANA Homo sapiens cell division cycle 27 (CDC27), mRNA
NM_001256	Homo sapiens CDC23 (cell division cycle 23, yeast, homolog) (CDC23), mRNA  Homo sapiens CDC23 (cell division cycle 23, yeast, homolog) (CDC23), mRNA
NM_004661	Homo sapiens CDC23 (cell division cycle 23, years, nomera, the Homo sapiens cyclin D-type binding-protein 1 (CCNDBP1), transcript variant 2,
NM_037370	TD 3.7.4
NM 012142	mRNA Homo sapiens cyclin D-type binding-protein 1 (CCNDBP1), transcript variant 1,
	mRNA
NM 019592	Homo sapiens ring finger protein 20 (RNF20), mRNA
NM 003386	ITama agriena zonadhesin (ZAN) mRNA
NM_001959	Homo sapiens eukaryotic translation elongation factor 1 beta 2 (EEF1B2),
1111_00120	Language of the second of the
NM 021121	Homo sapiens eukaryotic translation elongation factor I beta 2 (EEFIB2),
11112	the against various 2 mRNA
NM 006778	Home copiens ring finger protein 9 (RNF9), transcript variant 1, mRNA
NM 052828	Transcribe variance finger protein Q (R NF9). Transcribe variant 2, include
NM_007028	Homo sapiens tripartite motif-containing 31 (1RIM31), transcript variant 1,
NG 000010	Homo sapiens chorionic gonadotropin beta region (CGB@) on chromosome 19
NG 000019	Homo sapiens disrupted in renal carcinoma 1 (DIRC1), mRNA
NM 052952	Homo sapiens ribosomal protein L30 (RPL30), mRNA
NM_000989	Homo sapiens ribosomal protein L23 (RPL23), mRNA
NM_000978	Homo sapiens ribosomal protein L17 (RPL17), mRNA
NM_000985	Homo sapiens protocadherin 18 (PCDH18), mRNA
NM_019035	Homo sapiens nuclear RNA export factor 2 (NXF2), transcript variant 1, mRNA
NM_017809	Homo sapiens amnionless protein (AMN), mRNA
NM_030943	Homo sapiens animomess protein (14447), interest animomess protein (14447), interest variant 2, mRNA  Homo sapiens nuclear RNA export factor 2 (NXF2), transcript variant 2, mRNA
NM_022053	Homo sapiens 24-dehydrocholesterol reductase (DHCR24), mRNA
NM_014762	Homo sapiens 24-denydrocholesteror reduces (STAS2R14), mRNA  Homo sapiens taste receptor, type 2, member 14 (TAS2R14), mRNA
NM_023922	Homo sapiens taste receptor, type 2, member 17 (TAS2R10), mRNA  Homo sapiens taste receptor, type 2, member 10 (TAS2R10), mRNA
NM_023921	Homo sapiens taste receptor, type 2, member 13 (TAS2R13), mRNA  Homo sapiens taste receptor, type 2, member 13 (TAS2R13), mRNA
NM_023920	Homo sapiens taste receptor, type 2, member 7 (TAS2R7), mRNA  Homo sapiens taste receptor, type 2, member 7 (TAS2R7), mRNA
NM_023919	Homo sapiens taste receptor, type 2, member 7 (TAS2R8), mRNA  Homo sapiens taste receptor, type 2, member 8 (TAS2R8), mRNA
NM_023918	Homo sapiens taste receptor, type 2, member 9 (TAS2R9), mRNA  Homo sapiens taste receptor, type 2, member 9 (TAS2R9), mRNA
NM_023917	Homo sapiens taste receptor, type 2, memoer 5 (118287), nuclear gene  Homo sapiens mitochondrial ribosomal protein S14 (MRPS14), nuclear gene
NM_022100	Homo sapiens miliochondrial riocsomal protein BY (2.222 2.37)
	encoding mitochondrial protein, mRNA  Homo sapiens ATP-binding cassette, sub-family G (WHITE), member 4
NM_022169	Homo sapiens ATF-binding cassette, sub-taining G
	(ABCG4), mRNA  Homo sapiens WD repeat domain 6 (WDR6), transcript variant 1, mRNA
NM_018031	Homo sapiens c-myc binding protein (MYCBP), mRNA
NM_012333	Homo sapiens c-myc bilding protein (MTCBX), Med Homo sapiens hormonally upregulated Neu-associated kinase (HUNK), mRNA
NM_014586	Homo sapiens normanally upregulated recu-associated annual (
NM_014296	Homo sapiens calpain 7 (CAPN7), mRNA
NM 006615	
NM_005807	Homo sapiens proteoglycan 4, (megakaryocyte stimulating lactor, activities superficial zone protein, camptodactyly, arthropathy, coxa vara, pericarditis
}	superficial zone protein, campionactyly, artificipality, conditional value, periodically and the conditional value, periodically and the conditional value, periodically artificipality, conditional value, periodically artificipality artificipality, conditional value, and condition
	syndrome) (PRG4), mRNA
NM_004467	The state of the s
NM_003391	mDNI A
NM 002995	Homo sapiens small inducible cytokine subfamily C, member 1 (lymphotactin)

F	(SCYC1), mRNA
NM 002477	Homo sapiens myosin, light polypeptide 5, regulatory (MYL5), mRNA
NM 058253	Homo sapiens ribosomal protein S6 kinase, 52kD, polypeptide 1 (RPS6KC1),
11111_000000	mRNA
NM 000623	Homo sapiens bradykinin receptor B2 (BDKRB2), mRNA
NM 000424	Homo sapiens keratin 5 (epidermolysis bullosa simplex, Dowling-
_	Meara/Kobner/Weber-Cockayne types) (KRT5), mRNA
NM_002272	Homo sapiens keratin 4 (KRT4), mRNA
NM_057088	Homo sapiens keratin 3 (KRT3), mRNA
NM_006121	Homo sapiens keratin 1 (epidermolytic hyperkeratosis) (KRT1), mRNA
NM_057182	Homo sapiens cyclin E1 (CCNE1), transcript variant 2, mRNA
NM_001238	Homo sapiens cyclin E1 (CCNE1), transcript variant 1, mRNA
NM_054029	Homo sapiens chromosome 8 open reading frame 14 (C8orf14), mRNA
NM_054017	Homo sapiens chromosome 8 open reading frame 12 (C8orf12), mRNA
NM_052936	Homo sapiens AUT-like 2, cysteine endopeptidase (S. cerevisiae) (AUTL2), mRNA
NM_004926	Homo sapiens zinc finger protein 36, C3H type-like 1 (ZFP36L1), mRNA
NM_006887	Homo sapiens zinc finger protein 36, C3H type-like 2 (ZFP36L2), mRNA
NM 015355	Homo sapiens joined to JAZF1 (JJAZ1), mRNA
NM_005642	Homo sapiens TAF7 RNA polymerase II, TATA box binding protein (TBP)-
	associated factor, 55 kD (TAF7), mRNA
NM_032685	Homo sapiens hypothetical protein MGC13005 (MGC13005), mRNA
NM_032656	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 37 (DDX37), mRNA
NM 031919	Homo sapiens cystatin and DUF19 domain containing 1 (CSDUFD1), mRNA
NM 031475	Homo sapiens espin (ESPN), mRNA
NM_024101	Homo sapiens melanophilin (MLPH), mRNA
NM_002597	Homo sapiens phosducin (PDC), transcript variant Phd, mRNA
NM_021201	Homo sapiens membrane-spanning 4-domains, subfamily A, member 7 (MS4A7), mRNA
NM_020634	Homo sapiens growth differentiation factor 3 (GDF3), mRNA
NM_020185	Homo sapiens mitogen-activated protein kinase phosphatase x (MKPX), mRNA
NM_002897	Homo sapiens RNA binding motif, single stranded interacting protein 1 (RBMS1), transcript variant scr2, mRNA
NM_016839	Homo sapiens RNA binding motif, single stranded interacting protein 1
NM 016838	(RBMS1), transcript variant MSSP-2, mRNA  Homo sapiens RNA binding motif, single stranded interacting protein 1
14M_010036	(RBMS1), transcript variant MSSP-1, mRNA
NM 016837	Homo sapiens RNA binding motif, single stranded interacting protein 1
1411_010057	(RBMS1), transcript variant MSSP-3, mRNA
NM 016836	Homo sapiens RNA binding motif, single stranded interacting protein 1
71112_010000	(RBMS1), transcript variant YC1, mRNA
NM 016941	Homo sapiens delta-like 3 (Drosophila) (DLL3), mRNA
NM 016335	Homo sapiens proline dehydrogenase (oxidase) 1 (PRODH), mRNA
NM 014122	Homo sapiens PRO0245 protein (PRO0245), mRNA
NM 015344	Homo sapiens leptin receptor overlapping transcript-like 1 (LEPROTL1), mRNA
NM 014450	Homo sapiens SHP2 interacting transmembrane adaptor (SIT), mRNA
NM_007159	Homo sapiens sarcolemma associated protein (SLMAP), mRNA
NM_005974	Homo sapiens proline dehydrogenase (oxidase) 1 (PRODH), mRNA
NM_004974	Homo sapiens potassium voltage-gated channel, shaker-related subfamily, member 2 (KCNA2), mRNA
NM 003195	Homo sapiens transcription elongation factor A (SII), 2 (TCEA2), mRNA

	TO THE STATE OF TH
NM_001010	Homo sapiens ribosomal protein S6 (RPS6), mRNA
NM_000981	Homo sapiens ribosomal protein L19 (RPL19), mRNA
NM_003378	Homo sapiens VGF nerve growth factor inducible (VGF), mRNA  Homo sapiens VGF nerve growth factor inducible (VGF), mRNA
NM 001612	Homo sapiens acrosomal vesicle protein 1 (ACRV1), transcript variant 1, mRNA
NM 020115	Homo sapiens acrosomal vesicle protein 1 (ACRV1), transcript variant 11,
	mRNA mRNA
NM 020114	Homo sapiens acrosomal vesicle protein 1 (ACRV1), transcript variant 9, mRNA
NM 020113	Tr region protein   (Al. KVI), lialistipt variant 0, med 12
NM 020112	Tr
NM 020111	II was senione agreeomal vesicle protein I (ACRVI), transcript variant 0, matrix
NM 020110	Homo sapiens acrosomal vesicle protein 1 (ACRV1), transcript variant 10,
14141_020110	DATA
NM_020109	The serious coregonal vesicle protein 1 (ACRVI), transcript variant 5, filking
NM 020108	YV   A compromed vericle protein   (A(RVI), transcript variant 4, mid VI)
NM 020107	TY arrives coronal vesicle protein (ACRVI), transcript variant 5, inclvi
NM 020069	Homo saniens acrosomal vesicle protein I (ACRVI), transcript variant 2, micryi
NM 022909	TY and a contraction of (CENPH) MKNA
NM 021734	Homo sapiens solute carrier family 25 (mitochondrial deoxynucleotide carrier),
NWI_021/34	1 10 (CT C25 A 10) mPNA
ND ( 021250	Homo sapiens transmembrane protein 8 (five membrane-spanning domains)
NM_021259	(TMEMS) mPNA
ND ( 020120	Home conjens evidereductase UCPA (LOC56898), mRNA
NM_020139	Homo sapiens TAF9-like RNA polymerase II, TATA box binding protein
NM_015975	1 cmp m · · · · · · · · · · · · · · · · · ·
ND4 012271	Homo sapiens proprotein convertase subtilisin/kexin type 1 inhibitor (PCSK1N),
NM_013271	mRNA
ATA 6,000004	Homo saniens NAD(P)H dehydrogenase, quinone 2 (NQO2), mRNA
NM_000904	Homo sapiens NAD(P)H dehydrogenase, quinone 1 (NQO1), mRNA
NM_000903	TI was remiens contilin 1 (SORTI) mRNA
NM_002959	Homo sapiens G protein-coupled receptor kinase-interactor 2 (GIT2), transcript
NM_057170	
17.5 057160	Homo sapiens G protein-coupled receptor kinase-interactor 2 (GIT2), transcript
NM_057169	variant 1, mRNA
2 2 2 2 2 2 2 2 2	Ty
NM_057161	Homo sapiens testis intractitual includes processing from the sapiens collagen, type VI, alpha 3 (COL6A3), transcript variant 4, mRNA
NM_057167	11 4 VII olabo 2 (I VII 6 A 3) Transcribi Valialit 4, IIII VII
NM_057166	Homo sapiens collagen, type VI, alpha 3 (COL6A3), transcript variant 3, mRNA  Homo sapiens collagen, type VI, alpha 3 (COL6A3), transcript variant 2, mRNA
NM_057165	Homo sapiens collagen, type VI, alpha 3 (COL6A3), transcript variant 2, mRNA  Homo sapiens collagen, type VI, alpha 3 (COL6A3), transcript variant 2, mRNA
NM_057164	Homo sapiens Conagen, type v1, applies (COLD: 12),  Homo sapiens G protein-coupled receptor kinase-interactor 2 (GIT2), transcript
NM_014776	: 42 DNIA
	variant 3, mRNA  Homo sapiens collagen, type VI, alpha 3 (COL6A3), transcript variant 1, mRNA
NM_004369	Homo sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump),
NM_001183	Homo sapiens Al Fase, HT transporting, 1980501112 (ATDCC1) mDNA
	subunit 1 (ATP6S1), mRNA  Homo sapiens adenosine A2a receptor (ADORA2A), mRNA
NM_000675	Homo sapiens adenosine Aza receptor (ADOR 127)
NM_033027	Homo sapiens AXIN1 up-regulated (AXUD1), mRNA
NM_002539	Homo sapiens ornithine decarboxylase 1 (ODC1), mRNA  Homo sapiens phosphatidylinositol 4-kinase, catalytic, alpha polypeptide
NM_058004	Homo sapiens phosphaticylinositol 4-killase, catalytic, alpha posygrand
	(PIK4CA), transcript variant 2, mRNA
NM_000992	Homo sapiens ribosomal protein L29 (RPL29), mRNA
NM_000984	Homo sapiens ribosomal protein L23a (RPL23A), mRNA  Homo sapiens ribosomal protein L23a (RPL23A), mRNA
NM_001289	
NM 018648	

	RNPs) (NOLA3), mRNA
NM 021947	Homo sapiens serine racemase (SRR), mRNA
NM 016579	Homo sapiens 8D6 antigen (8D6A), mRNA
NM 006849	Homo sapiens protein disulfide isomerase, pancreatic (PDIP), mRNA
NM 002650	Homo sapiens phosphatidylinositol 4-kinase, catalytic, alpha polypeptide
14141_002030	(PIK4CA), transcript variant 1, mRNA
NM 000988	Homo sapiens ribosomal protein L27 (RPL27), mRNA
NM 000987	Homo sapiens ribosomal protein L26 (RPL26), mRNA
NM 000986	Homo sapiens ribosomal protein L24 (RPL24), mRNA
NM 031964	Homo sapiens keratin associated protein 17.1 (KAP17.1), mRNA
NM 000420	Homo sapiens Kell blood group (KEL), mRNA
NM_052841	Homo sapiens serine/threonine kinase 22C (spermiogenesis associated)
1111_052041	(STK22C), mRNA
NM_017647	Homo sapiens FtsJ homolog 3 (E. coli) (FTSJ3), mRNA
NM 001845	Homo sapiens collagen, type IV, alpha 1 (COL4A1), mRNA
NM 016508	Homo sapiens cyclin-dependent kinase-like 3 (CDKL3), mRNA
NM 001261	Homo sapiens cyclin-dependent kinase 9 (CDC2-related kinase) (CDK9), mRNA
NM_033131	Homo sapiens wingless-type MMTV integration site family, member 3A
	(WNT3A), mRNA
NM 030753	Homo sapiens wingless-type MMTV integration site family, member 3 (WNT3),
_	mRNA
NM 003396	Homo sapiens wingless-type MMTV integration site family, member 15
-	(WNT15), mRNA
NM_004626	Homo sapiens wingless-type MMTV integration site family, member 11
	(WNT11), mRNA
NM_057176	Homo sapiens barttin (BSND), mRNA
NM_012079	Homo sapiens diacylglycerol O-acyltransferase homolog 1 (mouse) (DGAT1),
NTM 005400	mRNA Homo sapiens SH2 domain-containing 3A (SH2D3A), mRNA
NM_005490 NM_032563	Homo sapiens shi domain-containing 5% (S11255%), met vi Homo sapiens epidermal differentiation complex protein like protein (LEP16),
NM_032363	mRNA
NM 014914	Homo sapiens centaurin, gamma 2 (CENTG2), mRNA
NM 014161	Homo sapiens mitochondrial ribosomal protein L18 (MRPL18), mRNA
NM 004895	Homo sapiens cold autoinflammatory syndrome 1 (CIAS1), mRNA
NM 000086	Homo sapiens ceroid-lipofuscinosis, neuronal 3, juvenile (Batten, Spielmeyer-
_	Vogt disease) (CLN3), mRNA
NM_033341	Homo sapiens baculoviral IAP repeat-containing 8 (BIRC8), mRNA
NM_054013	Homo sapiens mannosyl (alpha-1,3-)-glycoprotein beta-1,4-N-
	acetylglucosaminyltransferase, isoenzyme B (MGAT4B), transcript variant 2,
	mRNA
NM_000449	Homo sapiens regulatory factor X, 5 (influences HLA class II expression)
	(RFX5), mRNA
NM_054025	Homo sapiens beta-1,3-glucuronyltransferase 1 (glucuronosyltransferase P)
	(B3GAT1), transcript variant 2, mRNA
NM_002628	Homo sapiens profilin 2 (PFN2), transcript variant 2, mRNA
NM_053024	Homo sapiens profilin 2 (PFN2), transcript variant 1, mRNA
NM_003930	Homo sapiens src family associated phosphoprotein 2 (SCAP2), mRNA
NM_014018	Homo sapiens mitochondrial ribosomal protein S28 (MRPS28), nuclear gene
27.6.61.7071	encoding mitochondrial protein, mRNA
NM_015971	Homo sapiens mitochondrial ribosomal protein S7 (MRPS7), nuclear gene
DD 6 000 476	encoding mitochondrial protein, mRNA
NM_032476	Homo sapiens mitochondrial ribosomal protein S6 (MRPS6), nuclear gene

	encoding mitochondrial protein, mRNA
NM_018141	Homo sapiens mitochondrial ribosomal protein S10 (MRPS10), nuclear gene
1/1/1 010141	encoding mitochondrial protein, mRNA
ND4 014046	Homo sapiens mitochondrial ribosomal protein S18B (MRPS18B), nuclear gene
NM_014046	encoding mitochondrial protein, mRNA
ND 6 000512	Homo sapiens seryl-tRNA synthetase (SARS), mRNA
NM_006513	HOMO Sapiens Seryi-triva synthetase (SARO), mid 11
NM_021153	Homo sapiens cadherin 19, type 2 (CDH19), mRNA
NM_033664	Homo sapiens cadherin 11, type 2, OB-cadherin (osteoblast) (CDH11), transcript variant 2, mRNA
NM_001797	Homo sapiens cadherin 11, type 2, OB-cadherin (osteoblast) (CDH11), transcript variant 1, mRNA
NM_033381	Homo sapiens collagen, type IV, alpha 5 (Alport syndrome) (COL4A5), transcript variant 3, mRNA
NM_033380	Homo sapiens collagen, type IV, alpha 5 (Alport syndrome) (COL4A5), transcript variant 2, mRNA
NM_000495	Homo sapiens collagen, type IV, alpha 5 (Alport syndrome) (COL4A5), transcript variant 1, mRNA
NM 000092	Homo sapiens collagen, type IV, alpha 4 (COL4A4), mRNA
NM 033184	Homo sapiens keratin associated protein 2.4 (KAP2.4), mRNA
NM 032014	Homo sapiens mitochondrial ribosomal protein S24 (MRPS24), nuclear gene
11112_052011	encoding mitochondrial protein, mRNA
NM 001006	Homo sapiens ribosomal protein S3A (RPS3A), mRNA
NM 012411	Homo sapiens protein tyrosine phosphatase, non-receptor type 22 (lymphoid)
1111_012111	(PTPN22), transcript variant 2, mRNA
NM 015967	Homo sapiens protein tyrosine phosphatase, non-receptor type 22 (lymphoid)
1111_01050.	(PTPN22), transcript variant 1, mRNA
NM_006310	Homo sapiens aminopeptidase puromycin sensitive (NPEPPS), mRNA
NM_033335	Homo sapiens nuclear receptor subfamily 6, group A, member 1 (NR6A1),
1411_033333	transcript variant 3, mRNA
NM 033334	Homo sapiens nuclear receptor subfamily 6, group A, member 1 (NR6A1),
14141_055554	transcript variant 1, mRNA
NM 001489	Homo sapiens nuclear receptor subfamily 6, group A, member 1 (NR6A1),
14141_001403	transcript variant 2, mRNA
NM_001606	Homo sapiens ATP-binding cassette, sub-family A (ABC1), member 2 (ABCA2), mRNA
NM 002284	Homo sapiens keratin, hair, basic, 6 (monilethrix) (KRTHB6), mRNA
	Homo sapiens keratin, hair, basic, 5 (KRTHB5), mRNA
NM_002283	Homo sapiens keratin, hair, basic, 3 (KRTHB3), mRNA
NM_002282	Homo sapiens keratin, hair, basic, 2 (KRTHB2), mRNA
NM_033033	Homo sapiens keratin, hair, basic, 2 (KRTHB2), mRNA  Homo sapiens keratin, hair, basic, 1 (KRTHB1), mRNA
NM_002281	Homo sapiens keratin, hair, basic, 1 (KRTHB1), hikiya  Homo sapiens keratin, hair, basic, 4 (KRTHB4), mRNA
NM_033045	Homo sapiens keratin, nair, basic, 4 (KRTHB4), ilikiya  Homo sapiens ribosomal protein S7 (RPS7), mRNA
NM_001011	
NM_000980	Homo sapiens ribosomal protein L18a (RPL18A), mRNA
NM_000979	Homo sapiens ribosomal protein L18 (RPL18), mRNA
NM_000977	Homo sapiens ribosomal protein L13 (RPL13), transcript variant 1, mRNA
NM_033251	Homo sapiens ribosomal protein L13 (RPL13), transcript variant 2, mRNA
NM_000976	Homo sapiens ribosomal protein L12 (RPL12), mRNA
NM_000975	Homo sapiens ribosomal protein L11 (RPL11), mRNA
NM_000894	Homo sapiens luteinizing hormone beta polypeptide (LHB), mRNA
NM_005082	Homo sapiens zinc finger protein 147 (estrogen-responsive finger protein) (ZNF147), mRNA
NM 003549	Homo sapiens hyaluronoglucosaminidase 3 (HYAL3), mRNA

NM_033181	Homo sapiens cannabinoid receptor 1 (brain) (CNR1), transcript variant 3, mRNA
NG 000018	Homo sapiens genomic type I (acidic) hair keratin gene cluster (KRTHA.1@) on
149_00018	chromosome 17
NM 033151	Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 11
	(ABCC11), mRNA
NM_006998	Homo sapiens secretagogin (SECRET), mRNA
NM_006201	Homo sapiens PCTAIRE protein kinase 1 (PCTK1), transcript variant 1, mRNA
NM_033019	Homo sapiens PCTAIRE protein kinase 1 (PCTK1), transcript variant 3, mRNA
NM_033018	Homo sapiens PCTAIRE protein kinase 1 (PCTK1), transcript variant 2, mRNA
NG_000012	Homo sapiens genomic protocadherin gamma cluster (PCDHG@) on
	chromosome 5
NM_001023	Homo sapiens ribosomal protein S20 (RPS20), mRNA
NM_004451	Homo sapiens estrogen-related receptor alpha (ESRRA), mRNA
NM_005755	Homo sapiens Epstein-Barr virus induced gene 3 (EBI3), mRNA
NM_001015	Homo sapiens ribosomal protein S11 (RPS11), mRNA
NM_006923	Homo sapiens stromal cell-derived factor 2 (SDF2), mRNA
NM_000394	Homo sapiens crystallin, alpha A (CRYAA), mRNA
NM_003761	Homo sapiens vesicle-associated membrane protein 8 (endobrevin) (VAMP8),
	mRNA
NM_031958	Homo sapiens keratin associated protein 3.1 (KRTAP3.1), mRNA
NM_031957	Homo sapiens keratin associated protein 1.5 (KRTAP1.5), mRNA
NM_004776	Homo sapiens UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide
_	5 (B4GALT5), mRNA
NM_030587	Homo sapiens UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide
	2 (B4GALT2), transcript variant 1, mRNA
NM_003780	Homo sapiens UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide
	2 (B4GALT2), transcript variant 2, mRNA
NM_004391	Homo sapiens cytochrome P450, subfamily VIIIB (sterol 12-alpha-hydroxylase),
	polypeptide 1 (CYP8B1), mRNA
NM_000785	Homo sapiens cytochrome P450, subfamily XXVIIB (25-hydroxyvitamin D-1-
	alpha-hydroxylase), polypeptide 1 (CYP27B1), mitochondrial protein encoded
	by nuclear gene, mRNA
NM_031419	Homo sapiens molecule possessing ankyrin repeats induced by
	lipopolysaccharide (MAIL), homolog of mouse (MAIL), mRNA
NM_000961	Homo sapiens prostaglandin I2 (prostacyclin) synthase (PTGIS), mRNA
NM_003293	Homo sapiens tryptase, alpha (TPS1), mRNA
NM_016630	Homo sapiens acid cluster protein 33 (ACP33), mRNA
NM_014458	Homo sapiens Kelch motif containing protein (AB026190), mRNA
NM_007207	Homo sapiens dual specificity phosphatase 10 (DUSP10), mRNA
NM_030660	Homo sapiens Machado-Joseph disease (spinocerebellar ataxia 3,
	olivopontocerebellar ataxia 3, autosomal dominant, ataxin 3) (MJD), transcript
35.6	variant 2, mRNA
NM_022055	Homo sapiens potassium channel, subfamily K, member 12 (KCNK12), mRNA
NM_021175	Homo sapiens hepcidin antimicrobial peptide (HAMP), mRNA
NM_018666	Homo sapiens sarcoma antigen (SAGE), mRNA
NM_016532	Homo sapiens SKIP for skeletal muscle and kidney enriched inositol
	phosphatase (LOC51763), mRNA
NM_015987	Homo sapiens heme binding protein 1 (HEBP1), mRNA
NM_014079	Homo sapiens Kruppel-like factor 15 (KLF15), mRNA
NM_014759	Homo sapiens phytanoyl-CoA hydroxylase interacting protein (PHYHIP),
	mRNA

	interview 1 mDNA
NM_002590	Homo sapiens protocadherin 8 (PCDH8), transcript variant 1, mRNA
NM 004826	Homo sapiens endothelin converting enzyme-like 1 (ECEL1), mRNA
NM 004420	Homo sapiens dual specificity phosphatase 8 (DUSP8), mRNA
NM 001012	Homo sapiens ribosomal protein S8 (RPS8), mRNA
NM 002595	Homo saniens PCTAIRE protein kinase 2 (PCTK2), mRNA
NM 001395	Homo saniens dual specificity phosphatase 9 (DUSP9), mRNA
NM 003887	Homo sapiens development and differentiation enhancing factor 2 (DDEF2),
14141_002007	mRNA
NM 001446	Homo sapiens fatty acid binding protein 7, brain (FABP7), mRNA
NM 001259	Homo sapiens cyclin-dependent kinase 6 (CDK6), mRNA
	Homo sapiens cyclin D3 (CCND3), mRNA
NM_001760	Homo sapiens cyclin D2 (CCND2), mRNA
NM_001759	Homo sapiens cyclin A2 (CCNA2), mRNA
NM_001237	Homo sapiens cyclin A2 (CCNA2), inicial Homo sapiens dual specificity phosphatase 4 (DUSP4) transcript variant 2,
NM_057158	mPNA
NM_001394	Homo sapiens dual specificity phosphatase 4 (DUSP4), transcript variant 1, mRNA
NM_052988	Homo sapiens cyclin-dependent kinase (CDC2-like) 10 (CDK10), transcript
NM_052987	Homo sapiens cyclin-dependent kinase (CDC2-like) 10 (CDK10), transcript variant 2, mRNA
NM 057160	Homo sapiens artemin (ARTN), transcript variant 3, mRNA
NM 057091	Homo sapiens artemin (ARTN), transcript variant 2, mRNA
NM 057090	Homo sapiens artemin (ARTN), transcript variant 4, mRNA
NM 003976	Homo sapiens artemin (ARTN), transcript variant 1, mRNA
	Homo sapiens argininosuccinate synthetase (ASS), transcript variant 1, mRNA
NM_000050	Homo sapiens argininosuccinate synthetase (ASS), transcript variant 2, mRNA
NM_054012	Homo sapiens aquaporin 6, kidney specific (AQP6), transcript variant 2, mRNA
NM_053286	Homo sapiens aquaporin 6, kidney specific (AQP6), transcript variant 1, mRNA
NM_001652	Homo sapiens aquaporii o, kidney specific (NQ10), admissipe surface specific (NQ10), admissipe specific (NQ10), admissipe surface specific (NQ10), admissipe spec
NM_053032	mPNA
NM_053031	Homo sapiens myosin, light polypeptide kinase (MYLK), transcript variant 7, mRNA
NM_053030	Homo sapiens myosin, light polypeptide kinase (MYLK), transcript variant 5,
NM_053029	Homo sapiens myosin, light polypeptide kinase (MYLK), transcript variant 4,
NM_053028	Homo sapiens myosin, light polypeptide kinase (MYLK), transcript variant 3B,
NM_053027	Homo sapiens myosin, light polypeptide kinase (MYLK), transcript variant 3A, mRNA
NM_053026	Homo sapiens myosin, light polypeptide kinase (MYLK), transcript variant 2, mRNA
NM_053025	Homo sapiens myosin, light polypeptide kinase (MYLK), transcript variant 1, mRNA
NM_016497	Homo sapiens mitochondrial ribosomal protein 64 (MRP64), nuclear gene encoding mitochondrial protein, mRNA
NM_024026	Homo sapiens mitochondrial ribosomal protein 63 (MRP63), nuclear gene encoding mitochondrial protein, mRNA
NM_021821	Homo sapiens mitochondrial ribosomal protein S35 (MRPS35), nuclear gene encoding mitochondrial protein, mRNA
377 - 005065	Homo sapiens myosin, light polypeptide kinase (MYLK), transcript variant 6,
NM_005965	nomo sapiens myosm, ngm porypeptide kmase (MT LEC), transcript variant o,

	mRNA
NM 016640	Homo sapiens mitochondrial ribosomal protein S30 (MRPS30), mRNA
NM 053035	Homo sapiens mitochondrial ribosomal protein S33 (MRPS33), transcript variant
1111_055055	2, nuclear gene encoding mitochondrial protein, mRNA
NM 016071	Homo sapiens mitochondrial ribosomal protein S33 (MRPS33), transcript variant
	1, nuclear gene encoding mitochondrial protein, mRNA
NM 031901	Homo sapiens mitochondrial ribosomal protein S21 (MRPS21), transcript variant
-	1, nuclear gene encoding mitochondrial protein, mRNA
NM 018997	Homo sapiens mitochondrial ribosomal protein S21 (MRPS21), transcript variant
	2, nuclear gene encoding mitochondrial protein, mRNA
NM_033363	Homo sapiens mitochondrial ribosomal protein S12 (MRPS12), transcript variant
	3, nuclear gene encoding mitochondrial protein, mRNA
NM_033362	Homo sapiens mitochondrial ribosomal protein S12 (MRPS12), transcript variant
	2, nuclear gene encoding mitochondrial protein, mRNA
NM_021144	Homo sapiens PC4 and SFRS1 interacting protein 1 (PSIP1), mRNA
NM_052953	Homo sapiens hypothetical protein LRP15 (LRP15), mRNA
NM_033207	Homo sapiens transmembrane protein HTMP10 (HTMP10), mRNA
NM_030649	Homo sapiens centaurin, beta 5 (CENTB5), mRNA
NM_023936	Homo sapiens mitochondrial ribosomal protein S34 (MRPS34), nuclear gene
	encoding mitochondrial protein, mRNA
NM_021107	Homo sapiens mitochondrial ribosomal protein S12 (MRPS12), transcript variant
	1, nuclear gene encoding mitochondrial protein, mRNA
NM_014322	Homo sapiens opsin 3 (encephalopsin, panopsin) (OPN3), mRNA
NM_001260	Homo sapiens cyclin-dependent kinase 8 (CDK8), mRNA
NM_003674	Homo sapiens cyclin-dependent kinase (CDC2-like) 10 (CDK10), transcript
37.6.057004	variant 1, mRNA
NM_057094	Homo sapiens crystallin, beta A2 (CRYBA2), transcript variant 3, mRNA
NM_057093	Homo sapiens crystallin, beta A2 (CRYBA2), transcript variant 2, mRNA
NM_052984	Homo sapiens cyclin-dependent kinase 4 (CDK4), transcript variant 2, mRNA
NM_000075	Homo sapiens cyclin-dependent kinase 4 (CDK4), transcript variant 1, mRNA
NM_052827	Homo sapiens cyclin-dependent kinase 2 (CDK2), transcript variant 2, mRNA
NM_001798	Homo sapiens cyclin-dependent kinase 2 (CDK2), transcript variant 1, mRNA
NM_006522	Homo sapiens wingless-type MMTV integration site family, member 6 (WNT6), mRNA
NM_005430	Homo sapiens wingless-type MMTV integration site family, member 1 (WNT1), mRNA
NM_003394	Homo sapiens wingless-type MMTV integration site family, member 10B
	(WNT10B), mRNA
NM_025216	Homo sapiens wingless-type MMTV integration site family, member 10A (WNT10A), mRNA
NM_005370	Homo sapiens mel transforming oncogene (derived from cell line NK14)- RAB8 homolog (MEL), mRNA
NM_033100	Homo sapiens MT-protocadherin (KIAA1775), mRNA
NM 005086	Homo sapiens sarcospan (Kras oncogene-associated gene) (SSPN), mRNA
NM 003737	Homo sapiens protocadherin 16 (PCDH16), mRNA
NM 018153	Homo sapiens tumor endothelial marker 8 (TEM8), transcript variant 3, mRNA
NM 053034	Homo sapiens tumor endothelial marker 8 (TEM8), transcript variant 2, mRNA
NM_005929	Homo sapiens antigen p97 (melanoma associated) identified by monoclonal
	antibodies 133.2 and 96.5 (MFI2), transcript variant 1, mRNA
NM_033316	Homo sapiens antigen p97 (melanoma associated) identified by monoclonal
	antibodies 133.2 and 96.5 (MFI2), transcript variant 2, mRNA
NM_001002	Homo sapiens ribosomal protein, large, P0 (RPLP0), transcript variant 1, mRNA

	PO (DDI PO) transcript variant 2 mRNA
VM_053275	Homo sapiens ribosomal protein, large, P0 (RPLP0), transcript variant 2, mRNA
VM_054034	Homo sapiens fibronectin 1 (FN1), transcript variant 2, mRNA
NM_002026	Homo sapiens fibronectin 1 (FN1), transcript variant 1, mRNA
NM_004460	Homo sapiens fibroblast activation protein, alpha (FAP), mRNA  Homo sapiens fibroblast activation protein, alpha (FAP), mRNA
NM_000783	Homo sapiens rotochrome P450, subfamily XXVIA, polypeptide 1 (CYP26A1),
	transcript variant 1, mRNA
NM 057157	Homo sapiens cytochrome P450, subfamily XXVIA, polypeptide 1 (CYP26A1),
_	transcript variant 2. mRNA
NM 032211	Homo sapiens lysyl oxidase-like 4 (LOXL4), mRNA
NM_003395	Homo sapiens syst oxidase-inc + (1992-1993) Homo sapiens wingless-type MMTV integration site family, member 14
	(** P) T(D) 1 (A) D (A) (A)
NM 033101	Homo sapiens lectin, galactoside-binding, soluble, 12 (galectin 12) (LGALS12),
1,11,7_000	DNIA
NM_032611	Homo sapiens protein tyrosine phosphatase type IVA, member 3 (PTP4A3),
1111_032011	
NM_007079	Homo sapiens protein tyrosine phosphatase type IVA, member 3 (PTP4A3),
14141_007073	1
NM 032208	Home conjugate tumor endothelial marker 8 (TEM8), transcript variant 1, IRKNA
NM_014644	Homo sapiens phosphodiesterase 4D interacting protein (myomegalin)
NM_014044	(DDEADID) mDNA
NTM 006551	Homo sapiens lipophilin B (uteroglobin family member), prostatein-like
NM_006551	(LPHB), mRNA
NT ( 010000	Hama senions Ets I homolog 1 (F. coli) (FTSJ1), mRNA
NM_012280	Homo sapiens crystallin, beta A2 (CRYBA2), transcript variant 1, mRNA
NM_005209	Homo sapiens opioid growth factor receptor (OGFR), mRNA
NM_007346	Homo sapiens lipophilin A (uteroglobin family member) (LPHA), mRNA
NM_006552	Homo sapiens cell death-regulatory protein GRIM19 (GRIM19), mRNA
NM_015965	Homo sapiens mannosyl (alpha-1,3-)-glycoprotein beta-1,4-N-
NM_014275	Homo sapiens mannosyl (alpha-1,3-)-grycoprotein set 3, 1 acetylglucosaminyltransferase, isoenzyme B (MGAT4B), transcript variant 1,
	acetylglucosaminyltransletase, isochzynie B (MG11 12), danse i
	mRNA Homo sapiens carboxypeptidase B2 (plasma, carboxypeptidase U) (CPB2),
NM_001872	Homo sapiens carboxypeptidase bz (plastia, carboxypeptidase bz
	transcript variant 1, mRNA  Homo sapiens carboxypeptidase B2 (plasma, carboxypeptidase U) (CPB2),
NM_016413	Homo sapiens carboxypeptidase B2 (plasma, carboxypeptidase 3) (32 22)
	transcript variant 2, mRNA  Homo sapiens death associated protein 3 (DAP3), transcript variant 2, nuclear
NM_004632	Homo sapiens death associated protein 3 (DAF3), transcript variation,
	gene encoding mitochondrial protein, mRNA  Homo sapiens death associated protein 3 (DAP3), transcript variant 1, nuclear
NM_033657	Homo sapiens death associated protein 3 (DAF3), transcript variant 1, hadron
	gene encoding mitochondrial protein, mRNA
NM_001266	Homo sapiens carboxylesterase 1 (monocyte/macrophage serine esterase 1)
L	(CES1), mRNA
NM_004287	Homo sapiens golgi SNAP receptor complex member 2 (GOSR2), transcript
	variant A, mRNA  Homo sapiens golgi SNAP receptor complex member 2 (GOSR2), transcript
·	TI remains color CNAP recentor complex member 2 (UUSIX2), ualistript
NM_054022	Homo sapiens goigi Sivar receptor complex memory
NM_054022	variant B, mRNA
NM_054022 NM_002906	variant B, mRNA Homo sanjens radixin (RDX), mRNA
	Variant B, mRNA  Homo sapiens radixin (RDX), mRNA  Homo sapiens ribosomal protein, large P2 (RPLP2), mRNA
NM_002906 NM_001004	variant B, mRNA  Homo sapiens radixin (RDX), mRNA  Homo sapiens ribosomal protein, large P2 (RPLP2), mRNA  Homo sapiens ribosomal protein large, P1 (RPLP1), mRNA
NM_002906 NM_001004 NM_001003	variant B, mRNA  Homo sapiens radixin (RDX), mRNA  Homo sapiens ribosomal protein, large P2 (RPLP2), mRNA  Homo sapiens ribosomal protein, large, P1 (RPLP1), mRNA  Homo sapiens beta-1,3-glucuronyltransferase 1 (glucuronosyltransferase P)
NM_002906 NM_001004	variant B, mRNA  Homo sapiens radixin (RDX), mRNA  Homo sapiens ribosomal protein, large P2 (RPLP2), mRNA  Homo sapiens ribosomal protein, large, P1 (RPLP1), mRNA  Homo sapiens beta-1,3-glucuronyltransferase 1 (glucuronosyltransferase P)  (P3GAT1) transcript variant 1, mRNA
NM 002906 NM 001004 NM 001003 NM 018644	variant B, mRNA  Homo sapiens radixin (RDX), mRNA  Homo sapiens ribosomal protein, large P2 (RPLP2), mRNA  Homo sapiens ribosomal protein, large, P1 (RPLP1), mRNA  Homo sapiens beta-1,3-glucuronyltransferase 1 (glucuronosyltransferase P)  (B3GAT1), transcript variant 1, mRNA
NM_002906 NM_001004 NM_001003	variant B, mRNA  Homo sapiens radixin (RDX), mRNA  Homo sapiens ribosomal protein, large P2 (RPLP2), mRNA  Homo sapiens ribosomal protein, large, P1 (RPLP1), mRNA  Homo sapiens beta-1,3-glucuronyltransferase 1 (glucuronosyltransferase P)

NM_020439	Homo sapiens calcium/calmodulin-dependent protein kinase IG (CAMK1G), mRNA
NM_032158	Homo sapiens NOL1R2 protein (NOL1R2), mRNA
NM_022470	Homo sapiens p53 target zinc finger protein (WIG1), mRNA
NM_018044	Homo sapiens NOL1R protein (NOL1R), mRNA
NM_016262	Homo sapiens epsilon-tubulin (LOC51175), mRNA
NM_014239	Homo sapiens eukaryotic translation initiation factor 2B, subunit 2 (beta, 39kD) (EIF2B2), mRNA
NM_002308	Homo sapiens lectin, galactoside-binding, soluble, 9 (galectin 9) (LGALS9), transcript variant short, mRNA
NM_009587	Homo sapiens lectin, galactoside-binding, soluble, 9 (galectin 9) (LGALS9), transcript variant long, mRNA
NM_001187	Homo sapiens B melanoma antigen (BAGE), mRNA
NM_022162	Homo sapiens caspase recruitment domain family, member 15 (CARD15), mRNA
NM_014733	Homo sapiens endosome-associated FYVE-domain protein (ENDOFIN), mRNA
NM_013393	Homo sapiens FtsJ homolog 2 (E. coli) (FTSJ2), mRNA
NM_006440	Homo sapiens thioredoxin reductase beta (TR), mRNA
NM_005863	Homo sapiens neuroepithelial cell transforming gene 1 (NET1), mRNA
NM_002119	Homo sapiens major histocompatibility complex, class II, DO alpha (HLA-DOA), mRNA
NM_021908	Homo sapiens suppression of tumorigenicity 7 (ST7), transcript variant b, mRNA
NM_018412	Homo sapiens suppression of tumorigenicity 7 (ST7), transcript variant a, mRNA
NM_054020	Homo sapiens putative ion channel protein CATSPER2 (CATSPER2), mRNA
NM_053281	Homo sapiens dachshund homolog 2 (Drosophila) (DACH2), mRNA
NM_031439	Homo sapiens SOX7 transcription factor (SOX7), mRNA
NM_030796	Homo sapiens hypothetical protein DKFZp564K0822 (DKFZP564K0822), mRNA
NM_025117	Homo sapiens hypothetical protein FLJ11871 (FLJ11871), mRNA
NM_014893	Homo sapiens KIAA0951 protein (KIAA0951), mRNA
NM_000113	Homo sapiens dystonia 1, torsion (autosomal dominant; torsin A) (DYT1), mRNA
NM_053055	Homo sapiens C-terminal modulator protein (CTMP), mRNA
NM_021212	Homo sapiens HCF-binding transcription factor Zhangfei (ZF), mRNA
NM_007237	Homo sapiens SP140 nuclear body protein (SP140), mRNA
NM_006368	Homo sapiens cAMP responsive element binding protein 3 (luman) (CREB3), mRNA
NM_005759	Homo sapiens abl-interactor 12 (SH3-containing protein) (AIP-1), mRNA
NM_052966	Homo sapiens chromosome 1 open reading frame 24 (C1orf24), mRNA
NM_013247	Homo sapiens protease, serine, 25 (PRSS25), mRNA
NM_003017	Homo sapiens splicing factor, arginine/serine-rich 3 (SFRS3), mRNA
NM_006289	Homo sapiens talin 1 (TLN1), mRNA
NM_000970	Homo sapiens ribosomal protein L6 (RPL6), mRNA
NM_003973	Homo sapiens ribosomal protein L14 (RPL14), mRNA
NM 001361	Homo sapiens dihydroorotate dehydrogenase (DHODH), nuclear gene encoding
_	
	mitochondrial protein, mRNA
NM_021248	mitochondrial protein, mRNA Homo sapiens cadherin-like 22 (CDH22), mRNA
NM_021248 NM_033224	mitochondrial protein, mRNA  Homo sapiens cadherin-like 22 (CDH22), mRNA  Homo sapiens purine-rich element binding protein B (PURB), mRNA
NM_021248	mitochondrial protein, mRNA Homo sapiens cadherin-like 22 (CDH22), mRNA

	DATA
NM_013444	Homo sapiens ubiquilin 2 (UBQLN2), mRNA
NM 053067	Homo sapiens ubiquilin 1 (UBQLN1), transcript variant 2, mRNA
NM 013438	Homo sapiens ubiquilin 1 (UBQLN1), transcript variant 1, mRNA  Homo sapiens ubiquilin 1 (UBQLN1), transcript variant 1, mRNA
NM 032115	Homo sapiens ubiquim 1 (OBQEN1), transcript visiting protein (WEIKKN)  Homo sapiens potassium channel, subfamily K, member 16 (KCNK16), mRNA
NM_053284	Homo sapiens WAP, FS, Ig, KU, and NTR-containing protein (WFIKKN),
- 1	mDNA
NM 053278	Homo sapiens G protein-coupled receptor 102 (GPR102), mRNA
	TT MDNA
NM_032649	Homo sapiens glutamate carboxypeptidase-like protein 2 (CPGL2), ilitava
NM 053012	Homo sapiens hypothetical protein (LOC114137), mkNA
NM 003268	Homo sapiens toll-like receptor 5 (TLR5), mRNA
NM 053005	Homo sapiens HCCA2 protein (HCCA2), mRNA
NM 052889	Hame seniens CARD only protein (COP), mRNA
NM 024740	ri disputed in hinolar disorder 1 (DIBD1), MRNA
NM 015721	Home capiens gem (nuclear organelle) associated protein 4 (GEWHV4), filte (12)
NM 003730	Homo saniens ribonuclease 6 precursor (RNASEOPL), IIIKNA
NM 030916	II was senione to superfamily recentor LNIK (LNIK), IIIKINA
	Homo sapiens potassium channel, subfamily K, member 15 (TASK-5)
NM_022358	OKCNIV 15) mPNA
27.6.000576	Homo sapiens phosducin (PDC), transcript variant PhLOP1, mRNA
NM_022576	Tree : CIDItoin (CIDI ) mRNA
NM_018269	Homo sapiens SIPL protein (SIPL), interview Homo sapiens spastic paraplegia 3A (autosomal dominant) (SPG3A), mRNA
NM_015915	Homo sapiens G protein-coupled receptor 74 (GPR74), mRNA
NM_053036	Homo sapiens paralemmin 2 (PALM2), mRNA
NM_053016	Homo sapiens hypothetical protein (LOC114138), mRNA
NM_053057	Transition 1 (CEDT1) mRNA
NM_052838	Homo sapiens septin 1 (SEF 11), med 11  Homo sapiens solute carrier family 4, sodium bicarbonate transporter-like,
NM_032034	Homo sapiens solute carrier family 4, southin steams
	member 11 (SLC4A11), mRNA  Homo sapiens golgi phosphoprotein 5 (GOLPH5), mRNA  (TEM 20.A.) mRNA
NM_031899	Homo sapiens TBP-interacting protein (TIP120A), mRNA
NM_018448	Homo sapiens 1BP-interacting protein (1H 12015); Homo sapiens cell adhesion molecule-related/down-regulated by oncogenes
NM_016952	Homo sapiens cell adnesion molecule-related down regulation
	(CDON), mRNA  Homo sapiens mitochondrial ribosomal protein L53 (MRPL53), mRNA  Protein L53 (MRPL53), mRNA
NM_053050	Homo sapiens mitochondrial ribosomar protein 1955 (MGC14327), mRNA
NM_053045	Homo sapiens hypothetical protein MGC14327 (MGC14327), mRNA  Homo sapiens hypothetical protein MGC14327 (MGC14327), mRNA
NM_017680	Homo sapiens asporin (LRR class 1) (ASPN), mRNA
NM_003914	Homo sapiens cyclin A1 (CCNA1), mRNA
NM_032387	Homo sapiens cyclin XI (Column), Manual Homo sapiens protein kinase, lysine deficient 4 (PRKWNK4), mRNA
NM_019093	Homo sapiens UDP glycosyltransferase 1 family, polypeptide A3 (UGT1A3),
	mRNA 1.5 illumelymentide A9 (IIGT1A9)
NM_021027	Homo sapiens UDP glycosyltransferase 1 family, polypeptide A9 (UGT1A9),
_	mRNA 1.6 11 malamentida A.8 (IIGTI A.8)
NM 019076	Homo sapiens UDP glycosyltransferase 1 family, polypeptide A8 (UGT1A8),
_	mRNA 1 C 13 - 1 - 1 - 1 - 1 - 1 - 1 - 1 - 1 - 1
NM_000463	Homo sapiens UDP glycosyltransferase 1 family, polypeptide A1 (UGT1A1),
	mRNA
NM_016608	Homo sapiens ALEX1 protein (ALEX1), mRNA
NM 016607	II and comiens ALEX3 protein (ALEX3), mRNA
NM 014860	Homo saniens SPTF-associated factor 65 gamma (STAF65(gamma)), mixix
NM_014782	Try and armodillo repeat protein ALEX2 (ALEX2), IIIKINA
NM_001072	TOD 1 1/2 -C 1 tomaly notypeptide A01U(111/AU).
14141_001072	mRNA Homo sapiens GM2 ganglioside activator protein (GM2A), mRNA

NM_001975	Homo sapiens enolase 2, (gamma, neuronal) (ENO2), mRNA
NM_001428	Homo sapiens enolase 1, (alpha) (ENO1), mRNA
NM_052836	Homo sapiens cadherin related 23 (CDH23), transcript variant 2, mRNA
NM 022124	Homo sapiens cadherin related 23 (CDH23), transcript variant 1, mRNA
NM 004063	Homo sapiens cadherin 17, LI cadherin (liver-intestine) (CDH17), mRNA
NM 004062	Homo sapiens cadherin 16, KSP-cadherin (CDH16), mRNA
NM 004933	Homo sapiens cadherin 15, M-cadherin (myotubule) (CDH15), mRNA
NM 001257	Homo sapiens cadherin 13, H-cadherin (heart) (CDH13), mRNA
NM_052819	Homo sapiens caspase recruitment domain protein 14 (CARD14), transcript variant 2, mRNA
NM_024110	Homo sapiens caspase recruitment domain protein 14 (CARD14), transcript variant 1, mRNA
NM_032415	Homo sapiens caspase recruitment domain family, member 11 (CARD11), mRNA
NM 014466	Homo sapiens tektin 2 (testicular) (TEKT2), mRNA
NM_053006	Homo sapiens serine/threonine kinase 22B (spermiogenesis associated) (STK22B), mRNA
NM_012083	Homo sapiens frequently rearranged in advanced T-cell lymphomas 2 (FRAT2), mRNA
NM_006922	Homo sapiens sodium channel, voltage-gated, type III, alpha polypeptide (SCN3A), mRNA
NM_005347	Homo sapiens heat shock 70kD protein 5 (glucose-regulated protein, 78kD) (HSPA5), mRNA
NM 003777	Homo sapiens dynein, axonemal, heavy polypeptide 11 (DNAH11), mRNA
NM_013282	Homo sapiens ubiquitin-like, containing PHD and RING finger domains, 1 (UHRF1), mRNA
NM 020886	Homo sapiens ubiquitin specific protease 28 (USP28), mRNA
NM 020843	Homo sapiens zinc finger protein 291 (ZNF291), mRNA
NM 024529	Homo sapiens chromosome 1 open reading frame 28 (Clorf28), mRNA
NM 053003	Homo sapiens SIGLEC-like 1 (SIGLECL1), mRNA
NM 033329	Homo sapiens SIGLEC-like 1 (SIGLECL1), mRNA
NM 015101	Homo sapiens chromosome 1 open reading frame 17 (Clorf17), mRNA
NM 032551	Homo sapiens G protein-coupled receptor 54 (GPR54), mRNA
NM 031898	Homo sapiens tektin 3 (TEKT3), mRNA
NM 025191	Homo sapiens chromosome 1 open reading frame 22 (Clorf22), mRNA
NM 022755	Homo sapiens chromosome 9 open reading frame 12 (C9orf12), mRNA
NM 021104	Homo sapiens ribosomal protein L41 (RPL41), mRNA
NM 017847	Homo sapiens chromosome 1 open reading frame 27 (Clorf27), mRNA
	Homo sapiens chromosome 1 open reading frame 26 (Clorf26), mRNA
NM_017673	Homo sapiens mitochondrial CCA-adding tRNA-nucleotidyltransferase
NM_016000	•
NM_015989	(MtCCA), mRNA  Homo sapiens cysteine sulfinic acid decarboxylase-related protein 2 (CSAD), mRNA
NM_014654	Homo sapiens syndecan 3 (N-syndecan) (SDC3), mRNA
NM 014837	Homo sapiens chromosome 1 open reading frame 16 (Clorf16), mRNA
NM 007179	Homo sapiens insulin-like 6 (INSL6), mRNA
NM 005478	Homo sapiens insulin-like 5 (INSL5), mRNA
NM_053000	Homo sapiens TIGA1 (TIGA1), mRNA
NM_052940	Homo sapiens hypothetical protein MGC8974 (MGC8974), mRNA
NM_052830	Homo sapiens gamma-glutamyltransferase-like 3 (GGTL3), mRNA
NM_053002	Homo sapiens no opposite paired repeat protein (NOPAR), mRNA
NM_052998	Homo sapiens ornithine decarboxylase-like protein (ODC-p), mRNA

ND 6 050006	
NM_052996	Homo sapiens PR domain containing 7 (PRDM7), mRNA
NM_052995	Homo sapiens Usher syndrome 3A (USH3A), mRNA
NM_007110	Homo sapiens telomerase-associated protein 1 (TEP1), mRNA
NM_033656	Homo sapiens WD repeat domain 9 (WDR9), transcript variant 2, mRNA
NM_018963	Homo sapiens WD repeat domain 9 (WDR9), transcript variant 1, mRNA
NM_017818	Homo sapiens WD repeat domain 8 (WDR8), mRNA
NM_033662	Homo sapiens WD repeat domain 4 (WDR4), transcript variant 3, mRNA
NM_033661	Homo sapiens WD repeat domain 4 (WDR4), transcript variant 2, mRNA
NM_018669	Homo sapiens WD repeat domain 4 (WDR4), transcript variant 1, mRNA
NM_017883	Homo sapiens WD repeat domain 13 (WDR13), mRNA
NM_052837	Homo sapiens secretory carrier membrane protein 3 (SCAMP3), transcript variant 2, mRNA
NM_005698	Homo sapiens secretory carrier membrane protein 3 (SCAMP3), transcript variant 1, mRNA
NM_005697	Homo sapiens secretory carrier membrane protein 2 (SCAMP2), mRNA
NM_004866	Homo sapiens secretory carrier membrane protein 1 (SCAMP1), transcript variant 1, mRNA
NM_052822	Homo sapiens secretory carrier membrane protein 1 (SCAMP1), transcript variant 2, mRNA
NM_052811	Homo sapiens ret finger protein 2 (RFP2), transcript variant 2, mRNA
NM 005798	Homo sapiens ret finger protein 2 (RFP2), transcript variant 1, mRNA
NM 052817	Homo sapiens midline 2 (MID2), transcript variant 2, mRNA
NM 012216	Homo sapiens midline 2 (MID2), transcript variant 1, mRNA
NM 000798	Homo sapiens dopamine receptor D5 (DRD5), mRNA
NM 000794	Homo sapiens dopamine receptor D1 (DRD1), mRNA
NM 000796	Homo sapiens dopamine receptor D3 (DRD3), transcript variant a, mRNA
NM 033663	Homo sapiens dopamine receptor D3 (DRD3), transcript variant e, mRNA
NM 033660	Homo sapiens dopamine receptor D3 (DRD3), transcript variant d, mRNA
NM 033659	Homo sapiens dopamine receptor D3 (DRD3), transcript variant c, mRNA
NM 033658	Homo sapiens dopamine receptor D3 (DRD3), transcript variant b, mRNA
NM 004934	Homo sapiens cadherin 18, type 2 (CDH18), mRNA
NM 004061	Homo sapiens cadherin 12, type 2 (N-cadherin 2) (CDH12), mRNA
NM_030622	Homo sapiens cytochrome P450, subfamily IIS, polypeptide 1 (CYP2S1), mRNA
NM_052831	Homo sapiens dJ55C23.6 gene (dJ55C23.6), mRNA
NM_052816	Homo sapiens tripartite motif-containing 31 (TRIM31), transcript variant 2, mRNA
NM_052812	Homo sapiens tripartite motif-containing 15 (TRIM15), transcript variant 2, mRNA
NM_052955	Homo sapiens transglutaminase Z (TGM7), mRNA
NM_052957	Homo sapiens putative nuclear protein (NAAR1), mRNA
NM_052851	Homo sapiens similar to RhoGAP (GT650), mRNA
NM_033229	Homo sapiens tripartite motif-containing 15 (TRIM15), transcript variant 1, mRNA
NM_018103	Homo sapiens leucine-rich repeat-containing 5 (LRRC5), mRNA
NM_014879	Homo sapiens G protein-coupled receptor 105 (GPR105), mRNA
NM_000797	Homo sapiens dopamine receptor D4 (DRD4), mRNA
NM 006596	Homo sapiens polymerase (DNA directed), theta (POLQ), mRNA
NM_052972	Homo sapiens leucine-rich alpha-2-glycoprotein (LRG), mRNA
NM 052967	Homo sapiens mas-related G protein-coupled MRG (MRG), mRNA
NM 052963	Homo sapiens mitochondrial topoisomerase I (TOP1MT), mRNA
NM 052962	Homo sapiens class II cytokine receptor (IL22RA2), mRNA
	1

NM_052961	Homo sapiens solute carrier family 26, member 8 (SLC26A8), mRNA
NM_052958	Homo sapiens vestibule-1 protein (VEST1), mRNA
NM_052954	Homo sapiens cysteine and tyrosine-rich protein 1 (CYYR1), mRNA
NM_052949	Homo sapiens RAS guanyl releasing protein 4 (RASGRP4), mRNA
NM_052934	Homo sapiens solute carrier family 26, member 9 (SLC26A9), mRNA
NM_052933	Homo sapiens testis specific, 13 (TSGA13), mRNA
NM_052932	Homo sapiens pro-oncosis receptor inducing membrane injury gene (PORIMIN), mRNA
NM_052891	Homo sapiens peptidoglycan recognition protein-I-alpha precursor (PGLYRPIalpha), mRNA
NM_052888	Homo sapiens KIAA0563-related gene (LOC114659), mRNA
NM_052887	Homo sapiens Toll-interleukin 1 receptor (TIR) domain-containing adapter protein (TIRAP), mRNA
NM 052886	Homo sapiens mal, T-cell differentiation protein 2 (MAL2), mRNA
NM 052882	Homo sapiens zinc finger, imprinted 3 (ZIM3), mRNA
NM 052880	Homo sapiens hypothetical protein MGC17330 (MGC17330), mRNA
NM 052875	Homo sapiens hypothetical protein MGC10485 (MGC10485), mRNA
NM 052874	Homo sapiens syntaxin1B2 (STX1B2), mRNA
NM 052863	Homo sapiens putative cytokine high in normal-1 (HIN-1), mRNA
NM 052862	Homo sapiens hypothetical protein MGC21854 (MGC21854), mRNA
NM 052861	Homo sapiens hypothetical protein MGC21675 (MGC21675), mRNA
NM 052853	Homo sapiens hypothetical protein MGC20727 (MGC20727), mRNA
NM 052848	Homo sapiens hypothetical protein MGC20255 (MGC20255), mRNA
NM 052845	Homo sapiens hypothetical protein MGC20496 (MGC20496), mRNA
NM 052842	Homo sapiens BCL2-like 12 (proline rich) (BCL2L12), mRNA
NM 052818	Homo sapiens hypothetical gene CG018 (CG018), mRNA
NM 032514	Homo sapiens microtubule-associated protein 1 light chain 3 alpha
	(MAP1LC3A), mRNA
NM_022829	Homo sapiens solute carrier family 13 (sodium-dependent dicarboxylate transporter), member 3 (SLC13A3), mRNA
NM_018835	Homo sapiens olfactory receptor, family 1, subfamily K, member 1 (OR1K1), mRNA
NM_006750	Homo sapiens syntrophin, beta 2 (dystrophin-associated protein A1, 59kD, basic component 2) (SNTB2), mRNA
NM 033641	Homo sapiens collagen, type IV, alpha 6 (COL4A6), transcript variant B, mRNA
NM 001847	Homo sapiens collagen, type IV, alpha 6 (COL4A6), transcript variant A, mRNA
NM 004359	Homo sapiens cell division cycle 34 (CDC34), mRNA
NM_033493	Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 9, mRNA
NM_033492	Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 8, mRNA
NM_033491	Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 7, mRNA
NM_033490	Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1),
NM 033489	transcript variant 6, mRNA  Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1),
14141_033469	
NM 033488	transcript variant 5, mRNA  Home capiens cell division avale 2 like 1 (PITSI RE proteins) (CDC2L1)
	Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 4, mRNA
NM_033487	Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 3, mRNA
NM 033486	Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1),

	transcript variant 2, mRNA
NM_001787	Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1),
	transcript variant 1, mRNA
NM_005983	Homo sapiens S-phase kinase-associated protein 2 (p45) (SKP2), transcript
	variant 1, mRNA
NM_032637	Homo sapiens S-phase kinase-associated protein 2 (p45) (SKP2), transcript
_	variant 2, mRNA
NM_021968	Homo sapiens H4 histone family, member E (H4FE), mRNA
NM_002748	Homo sapiens mitogen-activated protein kinase 6 (MAPK6), mRNA
NM 003527	Homo sapiens H2B histone family, member N (H2BFN), mRNA
NM 001000	Homo sapiens ribosomal protein L39 (RPL39), mRNA
NM 000999	Homo sapiens ribosomal protein L38 (RPL38), mRNA
NM 000998	Homo sapiens ribosomal protein L37a (RPL37A), mRNA
NM 000997	Homo sapiens ribosomal protein L37 (RPL37), mRNA
NM 022054	Homo sapiens potassium channel, subfamily K, member 13 (KCNK13), mRNA
NM 021161	Homo sapiens potassium channel, subfamily K, member 10 (TREK-2)
_	(KCNK10), mRNA
NM 003944	Homo sapiens selenium binding protein 1 (SELENBP1), mRNA
NM 033649	Homo sapiens fibroblast growth factor 18 (FGF18), transcript variant 2, mRNA
NM 004114	Homo sapiens fibroblast growth factor 13 (FGF13), transcript variant 1A, mRNA
NM 033642	Homo sapiens fibroblast growth factor 13 (FGF13), transcript variant 1B, mRNA
NM 016279	Homo sapiens cadherin 9, type 2 (T1-cadherin) (CDH9), mRNA
NM 001796	Homo sapiens cadherin 8, type 2 (CDH8), mRNA
NM 031891	Homo sapiens cadherin 20, type 2 (CDH20), mRNA
NM 006727	Homo sapiens cadherin 10, type 2 (T2-cadherin) (CDH10), mRNA
NM 033671	Homo sapiens cyclin B3 (CCNB3), transcript variant 2, mRNA
NM 033670	Homo sapiens cyclin B3 (CCNB3), transcript variant 1, mRNA
NM 033379	Homo sapiens cell division cycle 2, G1 to S and G2 to M (CDC2), transcript
14141_055575	variant 2, mRNA
NM 001786	Homo sapiens cell division cycle 2, G1 to S and G2 to M (CDC2), transcript
1111_001700	variant 1, mRNA
NM_004361	Homo sapiens cadherin 7, type 2 (CDH7), transcript variant b, mRNA
NM 033646	Homo sapiens cadherin 7, type 2 (CDH7), transcript variant a, mRNA
NM 017734	Homo sapiens palmdelphin (PALMD), mRNA
NM 052832	Homo sapiens solute carrier family 26, member 7 (SLC26A7), mRNA
NM 018718	Homo sapiens testis specific, 14 (TSGA14), mRNA
NM 015935	Homo sapiens CGI-01 protein (CGI-01), mRNA
NM 033120	Homo sapiens naked cuticle homolog 2 (Drosophila) (NKD2), mRNA
NM 033031	Homo sapiens cyclin B3 (CCNB3), transcript variant 3, mRNA
NM 012068	Homo sapiens cyclin B3 (CCNB3), transcript variant 3, mRNA  Homo sapiens activating transcription factor 5 (ATF5), mRNA
	Homo sapiens CA11 (LOC56287), mRNA
NM_019617 NM_018398	Homo sapiens CATT (LOC36287), INRNA  Homo sapiens calcium channel, voltage-dependent, alpha 2/delta 3 subunit
14141_018338	
NR4 010210	(CACNA2D3), mRNA  Home senions transpil DNA phodehodicatorese (TDP1), mPNA
NM_018319	Homo sapiens tyrosyl-DNA phodphodiesterase (TDP1), mRNA
NM_014404	Homo sapiens calcium channel, voltage-dependent, gamma subunit 5
NTN # 01440#	(CACNG5), mRNA
NM_014405	Homo sapiens calcium channel, voltage-dependent, gamma subunit 4
27 6 010111	(CACNG4), mRNA
NM_012114	Homo sapiens caspase 14, apoptosis-related cysteine protease (CASP14), mRNA
NM_006985	Homo sapiens nuclear pore complex interacting protein (NPIP), mRNA
NM_006816	Homo sapiens chromosome 5 open reading frame 8 (C5orf8), mRNA
NM_006539	Homo sapiens calcium channel, voltage-dependent, gamma subunit 3

	(CACNG3), mRNA
NM 004347	Homo sapiens caspase 5, apoptosis-related cysteine protease (CASP5), mRNA
NM_003862	Homo sapiens fibroblast growth factor 18 (FGF18), transcript variant 1, mRNA
NM 020770	Homo sapiens cingulin (KIAA1319), mRNA
NM 030778	Homo sapiens hypothetical protein PRO1331 (PRO1331), mRNA
NM 004927	Homo sapiens mitochondrial ribosomal protein L49 (MRPL49), mRNA
NM 031962	
NM 031961	Homo sapiens keratin associated protein 9.3 (KRTAP9.3), mRNA
NM 033456	Homo sapiens keratin associated protein 9.2 (KRTAP9.2), mRNA
	Homo sapiens potassium channel, subfamily K, member 7 (KCNK7), transcript variant E, mRNA
NM_031854	Homo sapiens keratin associated protein 4.12 (KRTAP4.12), mRNA
NM_033455	Homo sapiens potassium channel, subfamily K, member 7 (KCNK7), transcript variant D, mRNA
NM_033348	Homo sapiens potassium channel, subfamily K, member 7 (KCNK7), transcript variant B, mRNA
NM_033347	Homo sapiens potassium channel, subfamily K, member 7 (KCNK7), transcript variant A, mRNA
NM 033191	Homo sapiens keratin associated protein 9.4 (KAP9.4), mRNA
NM 033061	Homo sapiens keratin associated protein 4.7 (KAP4.7), mRNA
NM 033188	Homo sapiens keratin associated protein 4.7 (KAP 4.7), IRNA  Homo sapiens keratin associated protein 4.5 (KAP4.5), mRNA
NM 033062	Homo sapiens keratin associated protein 4.3 (KAP4.3), mRNA  Homo sapiens keratin associated protein 4.2 (KAP4.2), mRNA
NM 033059	Homo sapiens keratin associated protein 4.12 (KAP4.14), mRNA
NM 033060	Homo sapiens keratin associated protein 4.14 (KAP 4.14), inkNA  Homo sapiens keratin associated protein 4.10 (KAP4.10), mRNA
NM 033643	Homo sapiens ribosomal protein L36 (RPL36), transcript variant 1, mRNA
NM 015414	Homo sapiens ribosomal protein L36 (RPL36), transcript variant 1, mRNA
NM 007209	Homo sapiens ribosomal protein L35 (RPL35), mRNA
NM 000996	Homo sapiens ribosomal protein L35 (RPL35A), mRNA
NM 033637	Homo sapiens beta-transducin repeat containing (BTRC), transcript variant 1,
144_055057	mRNA
NM 033345	Homo sapiens regulator of G-protein signalling 8 (RGS8), mRNA
NM 033543	Homo sapiens hypothetical protein R29124 1 (R29124 1), mRNA
NM 033186	Homo sapiens keratin associated protein 4.13 (KAP4.13), mRNA
NM_033050	Homo sapiens G protein-coupled receptor 91 (GPR91), mRNA
NM 032728	Homo sapiens hypothetical protein MGC12921 (MGC12921), mRNA
NM 032910	Homo sapiens hypothetical protein MGC14136 (MGC14136), mRNA
NM 032857	Homo sapiens mitochondrial ribosomal protein L56 (MRPL56), mRNA
NM 032640	Homo sapiens hypothetical protein MGC10526 (MGC10526), mRNA
NM 032560	Homo sapiens MSTP033 protein (MSTP033), mRNA
NM 032524	Homo sapiens keratin associated protein 4.4 (KRTAP4.4), mRNA
NM 032351	Homo sapiens mitochondrial ribosomal protein L45 (MRPL45), mRNA
NM 031963	Homo sapiens keratin associated protein 9.8 (KRTAP9.8), mRNA
NM_031432	Homo sapiens uridine-cytidine kinase 1 (UCK1), mRNA
NM_031289	Homo sapiens hypothetical protein MGC3146 (MGC3146), mRNA
NM 031269	Homo sapiens PRO1386 protein (PRO1386), mRNA
NM 030975	Homo sapiens keratin associated protein 9.9 (KRTAP9.9), mRNA
NM_030817	Homo sapiens hypothetical protein DKFZp434F0318 (DKFZP434F0318),
	mRNA
NM_017970	Homo sapiens hypothetical protein FLJ10008 (FLJ10008), mRNA
NM_024510	Homo sapiens hypothetical protein MGC4368 (MGC4368), mRNA
NM_024325	Homo sapiens hypothetical protein MGC10715 (MGC10715), mRNA
NM_023914	Homo sapiens G protein-coupled receptor 86 (GPR86), mRNA
NM_022915	Homo sapiens mitochondrial ribosomal protein L44 (MRPL44), mRNA

	TOTAL
NM_022469	Homo sapiens hypothetical protein FLJ21195 similar to protein related to DAC
	and cerberus (FLJ21195), mRNA
NM_022344	Homo sapiens protein kinase Njmu-R1 (NJMU-R1), mRNA
NM_002924	Homo sapiens regulator of G-protein signalling 7 (RGS7), mRNA
NM_020402	Homo sapiens cholinergic receptor, nicotinic, alpha polypeptide 10 (CHRNA10),
	mRNA
NM_015420	Homo sapiens DKFZP564O0463 protein (DKFZP564O0463), mRNA
NM_016355	Homo sapiens hqp0256 protein (LOC51202), mRNA
NM_020370	Homo sapiens G protein-coupled receptor 84 (GPR84), mRNA
NM_019016	Homo sapiens hypothetical protein (FLJ20261), mRNA
NM_017872	Homo sapiens hypothetical protein FLJ20546 (FLJ20546), mRNA
NM_018373	Homo sapiens hypothetical protein FLJ11271 (FLJ11271), mRNA
NM_018277	Homo sapiens hypothetical protein FLJ10932 (FLJ10932), mRNA
NM_018242	Homo sapiens hypothetical protein FLJ10847 (FLJ10847), mRNA
NM_016055	Homo sapiens mitochondrial ribosomal protein L48 (MRPL48), mRNA
NM_016468	Homo sapiens hypothetical protein (LOC51241), mRNA
NM_014099	Homo sapiens PRO1768 protein (PRO1768), mRNA
NM_014964	Homo sapiens KIAA1065 protein (KIAA1065), mRNA
NM_014859	Homo sapiens KIAA0672 gene product (KIAA0672), mRNA
NM_014174	Homo sapiens HSPC144 protein (HSPC144), mRNA
NM_014156	Homo sapiens DKFZP564O0463 protein (DKFZP564O0463), mRNA
NM_015544	Homo sapiens DKFZP564K1964 protein (DKFZP564K1964), mRNA
NM 015681	Homo sapiens B9 protein (B9), mRNA
NM_012301	Homo sapiens atrophin-1 interacting protein 1; activin receptor interacting
_	protein 1 (KIAA0705), mRNA
NM_006856	Homo sapiens activating transcription factor 7 (ATF7), mRNA
NM_005714	Homo sapiens potassium channel, subfamily K, member 7 (KCNK7), transcript
	variant C, mRNA
NM_005756	Homo sapiens G protein-coupled receptor 64 (GPR64), mRNA
NM_005267	Homo sapiens gap junction protein, alpha 8, 50kD (connexin 50) (GJA8), mRNA
NM_003457	Homo sapiens zinc finger protein 207 (ZNF207), mRNA
NM_003184	Homo sapiens TATA box binding protein (TBP)-associated factor, RNA
	polymerase II, B, 150kD (TAF2B), mRNA
NM_003079	Homo sapiens SWI/SNF related, matrix associated, actin dependent regulator of
	chromatin, subfamily e, member 1 (SMARCE1), mRNA
NM_002815	Homo sapiens proteasome (prosome, macropain) 26S subunit, non-ATPase, 11
	(PSMD11), mRNA
NM_002577	Homo sapiens p21 (CDKN1A)-activated kinase 2 (PAK2), mRNA
NM_003867	Homo sapiens fibroblast growth factor 17 (FGF17), mRNA
NM_003885	Homo sapiens cyclin-dependent kinase 5, regulatory subunit 1 (p35) (CDK5R1),
	mRNA
NM_003939	Homo sapiens beta-transducin repeat containing (BTRC), transcript variant 2,
	mRNA
NM_001208	Homo sapiens basic transcription factor 3, like 1 (BTF3L1), mRNA
NM_033500	Homo sapiens hexokinase 1 (HK1), transcript variant 5, nuclear gene encoding
	mitochondrial protein, mRNA
NM_033498	Homo sapiens hexokinase 1 (HK1), transcript variant 4, nuclear gene encoding
	mitochondrial protein, mRNA
NM_033497	Homo sapiens hexokinase 1 (HK1), transcript variant 3, nuclear gene encoding
	mitochondrial protein, mRNA
NM_033496	Homo sapiens hexokinase 1 (HK1), transcript variant 2, nuclear gene encoding
1	mitochondrial protein, mRNA

mRNA NM_033636 Homo sapiens SCAN domain-containing 2 (SCAND2), transcript variant 5, mRNA NM_033635 Homo sapiens SCAN domain-containing 2 (SCAND2), transcript variant 4, mRNA NM_033634 Homo sapiens SCAN domain-containing 2 (SCAND2), transcript variant 3, mRNA NM_033633 Homo sapiens SCAN domain-containing 2 (SCAND2), transcript variant 2, mRNA NM_033633 Homo sapiens SCAN domain-containing 2 (SCAND2), transcript variant 1, mRNA NM_033637 Homo sapiens SCAN domain-containing 2 (SCAND2), transcript variant 1, mRNA NM_033467 Homo sapiens PTEN induced putative kinase 1 (PINK1), mRNA NM_032409 Homo sapiens PTEN induced putative kinase 1 (PINK1), mRNA NM_032409 Homo sapiens PTEN induced putative kinase 1 (PINK1), mRNA NM_00129 Homo sapiens Ac-like transposable element (ALTE), mRNA NM_001192 Homo sapiens hexokinase 3 (white cell) (HK3), nuclear gene encoding mitochondrial protein, mRNA NM_00118 Homo sapiens hexokinase 1 (HK1), transcript variant 1, nuclear gene encoding mitochondrial protein, mRNA NM_004728 Homo sapiens bexokinase 1 (HK1), transcript variant 1, nuclear gene encoding mitochondrial protein, mRNA NM_004728 Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 21 (DDX21), mRNA NM_004728 Homo sapiens PAB 38, member RAS oncogene family (RAB38), mRNA NM_0162148 Homo sapiens scytokine receptor-like factor 2 (CRLF2), mRNA NM_01622 Homo sapiens schromosome 1 open reading frame 9 (Clorf9), mRNA NM_016428 Homo sapiens chromosome 1 open reading frame 9 (Clorf9), mRNA NM_018475 Homo sapiens chromosome 1 open reading frame 9 (Clorf9), mRNA NM_030340 Homo sapiens chromosome 1 open reading frame 15 (Clorf25), mRNA NM_030934 Homo sapiens chromosome 1 open reading frame 6 (CSorf6), mRNA NM_030934 Homo sapiens chromosome 1 open reading frame 6 (CSorf6), mRNA NM_0309350 Homo sapiens chromosome 5 open reading frame 6 (CSorf6), mRNA NM_016604 Homo sapiens chromosome 5 open reading frame 6 (CSorf6), mRNA NM_035508 Homo sapiens chromosome 5 open reading frame 6 (CSorf6), mRNA NM_035507 Homo sapiens chromosome 6 open reading	NM 033640	Homo sapiens SCAN domain-containing 2 (SCAND2), transcript variant 6,
mRNA NM_033635 Homo sapiens SCAN domain-containing 2 (SCAND2), transcript variant 4, mRNA NM_033634 Homo sapiens SCAN domain-containing 2 (SCAND2), transcript variant 3, mRNA NM_033633 Homo sapiens SCAN domain-containing 2 (SCAND2), transcript variant 2, mRNA NM_022050 Homo sapiens SCAN domain-containing 2 (SCAND2), transcript variant 1, mRNA NM_033637 Homo sapiens PCAN domain-containing 2 (SCAND2), transcript variant 1, mRNA NM_033467 Homo sapiens membrane metallo-endopeptidase-like 2 (MMEL2), mRNA NM_032409 Homo sapiens breast cell glutaminase (GA), mRNA NM_0013267 Homo sapiens breast cell glutaminase (GA), mRNA NM_004729 Homo sapiens Ac-like transposable element (ALTE), mRNA NM_004192 Homo sapiens Ac-like transposable element (ALTE), mRNA NM_00115 Homo sapiens hexokinase 3 (white cell) (HK3), nuclear gene encoding mitochondrial protein, mRNA NM_00188 Homo sapiens hexokinase 1 (HK1), transcript variant 1, nuclear gene encoding mitochondrial protein, mRNA NM_004728 Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 21 (DDX21), mRNA NM_02148 Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 21 (DDX21), mRNA NM_02337 Homo sapiens NRSH protein (NESH), mRNA NM_016228 Homo sapiens NRSH protein (NESH), mRNA NM_016228 Homo sapiens chromosome 1 open reading frame 9 (Clorf9), mRNA NM_016248 Homo sapiens chromosome 1 open reading frame 9 (Clorf9), mRNA NM_016248 Homo sapiens chromosome 1 open reading frame 9 (Clorf9), mRNA NM_016428 Homo sapiens chromosome 1 open reading frame 25 (Clorf25), mRNA NM_030349 Homo sapiens chromosome 1 open reading frame 14 (Clorf14), mRNA NM_030934 Homo sapiens chromosome 1 open reading frame 15 (Clorf25), mRNA NM_030934 Homo sapiens chromosome 1 open reading frame 16 (C50rf5), mRNA NM_03060 Homo sapiens chromosome 5 open reading frame 6 (C50rf6), mRNA NM_01660 Homo sapiens chromosome 5 open reading frame 6 (C50rf6), mRNA NM_01660 Homo sapiens chromosome 5 open reading frame 7 (C50rf7), mRNA NM_01660 Homo sapiens chromosome 5 open reading frame 6 (C50rf6), mRNA NM_01660 Homo		mRNA
mRNA  NM_033634 Homo sapiens SCAN domain-containing 2 (SCAND2), transcript variant 3, mRNA  NM_033633 Homo sapiens SCAN domain-containing 2 (SCAND2), transcript variant 2, mRNA  NM_022050 Homo sapiens SCAN domain-containing 2 (SCAND2), transcript variant 1, mRNA  NM_033467 Homo sapiens membrane metallo-endopeptidase-like 2 (MMEL2), mRNA  NM_032409 Homo sapiens PTEN induced putative kinase 1 (PINK1), mRNA  NM_013267 Homo sapiens breast cell glutaminase (GA), mRNA  NM_004729 Homo sapiens Ac-like transposable element (ALTE), mRNA  NM_004729 Homo sapiens sectylserotonin O-methyltransferase-like (ASMTL), mRNA  NM_001151 Homo sapiens hexokinase 3 (white cell) (HK3), nuclear gene encoding mitochondrial protein, mRNA  NM_000188 Homo sapiens hexokinase 1 (HK1), transcript variant 1, nuclear gene encoding mitochondrial protein, mRNA  NM_04728 Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 21 (DDX21), mRNA  NM_022337 Homo sapiens RAB38, member RAS oncogene family (RAB38), mRNA  NM_016428 Homo sapiens NESH protein (NESH), mRNA  NM_016428 Homo sapiens Shromosome 1 open reading frame 9 (Clorf9), mRNA  NM_016428 Homo sapiens chromosome 1 open reading frame 9 (Clorf9), mRNA  NM_016428 Homo sapiens chromosome 1 open reading frame 9 (Clorf9), mRNA  NM_016428 Homo sapiens chromosome 1 open reading frame 9 (Clorf9), mRNA  NM_010604 Homo sapiens chromosome 1 open reading frame 25 (Clorf25), mRNA  NM_010605 Homo sapiens chromosome 1 open reading frame 14 (Clorf14), mRNA  NM_030934 Homo sapiens chromosome 1 open reading frame 13 (Clorf13), mRNA  NM_016604 Homo sapiens chromosome 5 open reading frame 6 (CSorf6), mRNA  NM_016605 Homo sapiens chromosome 5 open reading frame 7 (CSorf7), mRNA  NM_016604 Homo sapiens chromosome 5 open reading frame 6 (CSorf6), mRNA  NM_016605 Homo sapiens chromosome 5 open reading frame 6 (CSorf6), mRNA  NM_016604 Homo sapiens chromosome 5 open reading frame 6 (CSorf6), mRNA  NM_016605 Homo sapiens chromosome 5 open reading frame 6 (CSorf6), mRNA  NM_016606 Homo sapiens chromosome 5 open reading	NM_033636	
NM_033634   Homo sapiens SCAN domain-containing 2 (SCAND2), transcript variant 3, mRNA   mRNA   Homo sapiens SCAN domain-containing 2 (SCAND2), transcript variant 2, mRNA   Homo sapiens SCAN domain-containing 2 (SCAND2), transcript variant 1, mRNA   MM_033467   Homo sapiens membrane metallo-endopeptidase-like 2 (MMEL2), mRNA   MM_032409   Homo sapiens PTEN induced putative kinase 1 (PINK1), mRNA   NM_032409   Homo sapiens seast cell glutaminase (GA), mRNA   NM_032409   Homo sapiens seast cell glutaminase (GA), mRNA   NM_0404729   Homo sapiens acetylserotonin O-methyltransferase-like (ASMTL), mRNA   NM_004192   Homo sapiens hexokinase 3 (white cell) (HK3), nuclear gene encoding mitochondrial protein, mRNA   NM_000185   Homo sapiens hexokinase 1 (HK1), transcript variant 1, nuclear gene encoding mitochondrial protein, mRNA   NM_004728   Homo sapiens bexokinase 1 (HK1), transcript variant 1, nuclear gene encoding mitochondrial protein, mRNA   NM_004728   Homo sapiens cytokine receptor-like factor 2 (CRLF2), mRNA   NM_016428   Homo sapiens RAB38, member RAS oncogene family (RAB38), mRNA   NM_016428   Homo sapiens NESH protein (NESH), mRNA   NM_016427   Homo sapiens chromosome 1 open reading frame 9 (C1orf9), mRNA   NM_014283   Homo sapiens chromosome 1 open reading frame 9 (C1orf9), mRNA   NM_014284   Homo sapiens chromosome 1 open reading frame 9 (C1orf9), mRNA   NM_016404   Homo sapiens chromosome 1 open reading frame 25 (C1orf25), mRNA   NM_016604   Homo sapiens chromosome 1 open reading frame 13 (C1orf14), mRNA   NM_016605   Homo sapiens chromosome 5 open reading frame 6 (C5orf6), mRNA   NM_016605   Homo sapiens chromosome 5 open reading frame 7 (C5orf7), mRNA   NM_016605   Homo sapiens chromosome 5 open reading frame 6 (C5orf6), mRNA   NM_016605   Homo sapiens chromosome 5 open reading frame 7 (C5orf7), mRNA   NM_016605   Homo sapiens chromosome 5 open reading frame 7 (C5orf7), mRNA   NM_016605   Homo sapiens chromosome 5 open reading frame 6 (C5orf6), mRNA   NM_016605   Homo sapiens chromosome 6 open readin	NM_033635	
NM_023633	NM_033634	Homo sapiens SCAN domain-containing 2 (SCAND2), transcript variant 3,
NM_022050   Homo sapiens SCAN domain-containing 2 (SCAND2), transcript variant 1, mRNA   NM_033467   Homo sapiens membrane metallo-endopeptidase-like 2 (MMEL2), mRNA   NM_032409   Homo sapiens breast cell glutaminase (GA), mRNA   NM_041267   Homo sapiens breast cell glutaminase (GA), mRNA   NM_04192   Homo sapiens Ac-like transposable element (ALTE), mRNA   NM_04192   Homo sapiens acetylserotomin O-methyltransferase-like (ASMTL), mRNA   NM_04192   Homo sapiens hexokinase 3 (white cell) (HK3), nuclear gene encoding mitochondrial protein, mRNA   Homo sapiens hexokinase 1 (HK1), transcript variant 1, nuclear gene encoding mitochondrial protein, mRNA   Homo sapiens beach (Asp-Glu-Ala-Asp/His) box polypeptide 21 (DDX21), mRNA   NM_04728   Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 21 (DDX21), mRNA   NM_04728   Homo sapiens cytokine receptor-like factor 2 (CRLF2), mRNA   NM_04237   Homo sapiens RAB38, member RAS oncogene family (RAB38), mRNA   NM_04281   Homo sapiens bresh protein (NESH), mRNA   NM_04282   Homo sapiens chromosome 1 open reading frame 9 (Clorf9), mRNA   NM_04283   Homo sapiens chromosome 1 open reading frame 9 (Clorf9), mRNA   NM_04283   Homo sapiens pamma-tubulin complex component (GCP6), mRNA   NM_04461   Homo sapiens pamma-tubulin complex component (GCP6), mRNA   NM_0403093   Homo sapiens chromosome 1 open reading frame 25 (Clorf25), mRNA   NM_040604   Homo sapiens chromosome 1 open reading frame 13 (Clorf13), mRNA   NM_040605   Homo sapiens chromosome 5 open reading frame 14 (Clorf14), mRNA   NM_040603   Homo sapiens chromosome 5 open reading frame 6 (C5orf6), mRNA   NM_040603   Homo sapiens chromosome 5 open reading frame 6 (C5orf6), mRNA   NM_040603   Homo sapiens chromosome 5 open reading frame 6 (C5orf6), mRNA   NM_040603   Homo sapiens chromosome 5 open reading frame 6 (C5orf6), mRNA   NM_040604   Homo sapiens chromosome 6 open reading frame 7 (C5orf7), mRNA   NM_040605   Homo sapiens chromosome 6 open reading frame 7 (C5orf7), mRNA   NM_040606   Homo sapiens chromosome 6 ope	NM_033633	Homo sapiens SCAN domain-containing 2 (SCAND2), transcript variant 2,
NM_033467         Homo sapiens membrane metallo-endopeptidase-like 2 (MMEL2), mRNA           NM_032409         Homo sapiens PTEN induced putative kinase 1 (PINK1), mRNA           NM_013267         Homo sapiens breast cell glutaminase (GA), mRNA           NM_004729         Homo sapiens Ac-like transposable element (ALTE), mRNA           NM_004192         Homo sapiens hexokinase 3 (white cell) (HK3), nuclear gene encoding mitochondrial protein, mRNA           NM_002115         Homo sapiens hexokinase 1 (HK1), transcript variant 1, nuclear gene encoding mitochondrial protein, mRNA           NM_000188         Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 21 (DDX21), mRNA           NM_004728         Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 21 (DDX21), mRNA           NM_022148         Homo sapiens RAB38, member RAS oncogene family (RAB38), mRNA           NM_016428         Homo sapiens RAB38, member RAS oncogene family (RAB38), mRNA           NM_016427         Homo sapiens bexnomosome 1 open reading frame 9 (Clorf9), mRNA           NM_016227         Homo sapiens chromosome 1 open reading frame 9 (Clorf9), mRNA           NM_018475         Homo sapiens TPA regulated locus (TPARL), mRNA           NM_030934         Homo sapiens chromosome 1 open reading frame 25 (Clorf25), mRNA           NM_030934         Homo sapiens chromosome 1 open reading frame 14 (Clorf14), mRNA           NM_030604         Homo sapiens chromosome 5 open reading frame 6 (C5or	NM_022050	Homo sapiens SCAN domain-containing 2 (SCAND2), transcript variant 1,
NM032409Homo sapiens PTEN induced putative kinase 1 (PINK1), mRNANM013267Homo sapiens breast cell glutaminase (GA), mRNANM004729Homo sapiens Acc-like transposable element (ALTE), mRNANM004192Homo sapiens acetylserotonin O-methyltransferase-like (ASMTL), mRNANM002115Homo sapiens hexokinase 3 (white cell) (HK3), nuclear gene encoding mitochondrial protein, mRNANM000188Homo sapiens hexokinase 1 (HK1), transcript variant 1, nuclear gene encoding mitochondrial protein, mRNANM004728Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 21 (DDX21), mRNANM022148Homo sapiens cytokine receptor-like factor 2 (CRLF2), mRNANM022337Homo sapiens RAB38, member RAS oncogene family (RAB38), mRNANM016428Homo sapiens bromosome 1 open reading frame 9 (Clorf9), mRNANM014283Homo sapiens chromosome 1 open reading frame 9 (Clorf9), mRNANM014284Homo sapiens chromosome 1 open reading frame 9 (Clorf9), mRNANM018475Homo sapiens gamma-tubulin complex component (GCP6), mRNANM030934Homo sapiens chromosome 1 open reading frame 25 (Clorf25), mRNANM030934Homo sapiens chromosome 1 open reading frame 14 (Clorf14), mRNANM030769Homo sapiens chromosome 5 open reading frame 7 (CSorf7), mRNANM016604Homo sapiens chromosome 5 open reading frame 6 (CSorf6), mRNANM016605Homo sapiens chromosome 5 open reading frame 21 (Clorf121), mRNANM016606Homo sapiens chromosome 1 open reading frame	NM 033467	<u></u>
NM013267Homo sapiens breast cell glutaminase (GA), mRNANM004729Homo sapiens Ac-like transposable element (ALTE), mRNANM004192Homo sapiens acetylserotonin O-methyltransferase-like (ASMTL), mRNANM002115Homo sapiens hexokinase 3 (white cell) (HK3), nuclear gene encoding mitochondrial protein, mRNANM000188Homo sapiens hexokinase 1 (HK1), transcript variant 1, nuclear gene encoding mitochondrial protein, mRNANM004728Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 21 (DDX21), mRNANM002148Homo sapiens Cytokine receptor-like factor 2 (CRLF2), mRNANM022148Homo sapiens RAB38, member RAS oncogene family (RAB38), mRNANM016228Homo sapiens SESH protein (NESH), mRNANM016227Homo sapiens chromosome 1 open reading frame 9 (Clorf9), mRNANM018475Homo sapiens chromosome 1 open reading frame 9 (Clorf9), mRNANM018475Homo sapiens gamma-tubulin complex component (GCP6), mRNANM030934Homo sapiens chromosome 1 open reading frame 25 (Clorf25), mRNANM030933Homo sapiens chromosome 1 open reading frame 25 (Clorf25), mRNANM030933Homo sapiens chromosome 1 open reading frame 13 (Clorf13), mRNANM016604Homo sapiens chromosome 5 open reading frame 6 (CSorf6), mRNANM016605Homo sapiens chromosome 5 open reading frame 6 (CSorf6), mRNANM0164144Homo sapiens chromosome 5 open reading frame 21 (Cl1orf21), mRNANM003508Homo sapiens glucokinase (hexokinase 4, maturity onset diabe		
NM004729Homo sapiens Ac-like transposable element (ALTE), mRNANM004192Homo sapiens acetylserotonin O-methyltransferase-like (ASMTL), mRNANM002115Homo sapiens hexokinase 3 (white cell) (HK3), nuclear gene encoding mitochondrial protein, mRNANM000188Homo sapiens hexokinase 1 (HK1), transcript variant 1, nuclear gene encoding mitochondrial protein, mRNANM004728Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 21 (DDX21), mRNANM022148Homo sapiens cytokine receptor-like factor 2 (CRLF2), mRNANM022337Homo sapiens RAB38, member RAS oncogene family (RAB38), mRNANM016428Homo sapiens NESH protein (NESH), mRNANM016227Homo sapiens chromosome 1 open reading frame 9 (C1orf9), mRNANM014283Homo sapiens chromosome 1 open reading frame 9 (C1orf9), mRNANM018475Homo sapiens gamma-tubulin complex component (GCP6), mRNANM030934Homo sapiens chromosome 1 open reading frame 25 (C1orf25), mRNANM030933Homo sapiens chromosome 1 open reading frame 13 (C1orf13), mRNANM030694Homo sapiens chromosome 5 open reading frame 7 (C5orf7), mRNANM016604Homo sapiens chromosome 5 open reading frame 5 (C5orf5), mRNANM016605Homo sapiens chromosome 5 open reading frame 5 (C5orf5), mRNANM016604Homo sapiens chromosome 5 open reading frame 5 (C5orf5), mRNANM01603Homo sapiens chromosome 5 open reading frame 5 (C5orf5), mRNANM01604Homo sapiens chromosome 5 open reading frame 5 (C5orf5		
NM_004192Homo sapiens acetylserotonin O-methyltransferase-like (ASMTL), mRNANM_002115Homo sapiens hexokinase 3 (white cell) (HK3), nuclear gene encoding mitochondrial protein, mRNANM_000188Homo sapiens hexokinase 1 (HK1), transcript variant 1, nuclear gene encoding mitochondrial protein, mRNANM_004728Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 21 (DDX21), mRNANM_022148Homo sapiens cytokine receptor-like factor 2 (CRLF2), mRNANM_022337Homo sapiens RAB38, member RAS oncogene family (RAB38), mRNANM_016428Homo sapiens NESH protein (NESH), mRNANM_016227Homo sapiens chromosome 1 open reading frame 9 (Clorf9), mRNANM_018475Homo sapiens TPA regulated locus (TPARL), mRNANM_020461Homo sapiens chromosome 1 open reading frame 25 (Clorf25), mRNANM_030934Homo sapiens chromosome 1 open reading frame 25 (Clorf25), mRNANM_030933Homo sapiens chromosome 1 open reading frame 14 (Clorf14), mRNANM_030604Homo sapiens chromosome 5 open reading frame 13 (Clorf13), mRNANM_016605Homo sapiens chromosome 5 open reading frame 6 (C5orf6), mRNANM_016604Homo sapiens chromosome 5 open reading frame 5 (C5orf5), mRNANM_016605Homo sapiens chromosome 5 open reading frame 21 (Cl1orf21), mRNANM_033508Homo sapiens chromosome 10 open reading frame 21 (Cl1orf21), mRNANM_033507Homo sapiens chromosome 5 open reading frame 5 (C5orf5), mRNANM_033508Homo sapiens chromosome 6 open reading frame 5 (C5orf5), mRNANM_033507Homo sapiens chromosome 6 open reading frame 5 (C5orf5), mRNANM_033508Hom		
NM_002115  Homo sapiens hexokinase 3 (white cell) (HK3), nuclear gene encoding mitochondrial protein, mRNA  NM_000188  Homo sapiens hexokinase 1 (HK1), transcript variant 1, nuclear gene encoding mitochondrial protein, mRNA  NM_004728  Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 21 (DDX21), mRNA  NM_022148  Homo sapiens cytokine receptor-like factor 2 (CRLF2), mRNA  NM_022337  Homo sapiens RAB38, member RAS oncogene family (RAB38), mRNA  NM_016428  Homo sapiens NESH protein (NESH), mRNA  NM_016227  Homo sapiens chromosome 1 open reading frame 9 (Clorf9), mRNA  NM_014283  Homo sapiens chromosome 1 open reading frame 9 (Clorf9), mRNA  NM_018475  Homo sapiens TPA regulated locus (TPARL), mRNA  NM_030934  Homo sapiens chromosome 1 open reading frame 25 (Clorf25), mRNA  NM_030933  Homo sapiens chromosome 1 open reading frame 25 (Clorf25), mRNA  NM_030934  Homo sapiens chromosome 1 open reading frame 13 (Clorf13), mRNA  NM_030769  Homo sapiens chromosome 1 open reading frame 13 (Clorf13), mRNA  NM_016604  Homo sapiens chromosome 5 open reading frame 6 (CSorf6), mRNA  NM_016605  Homo sapiens chromosome 5 open reading frame 6 (CSorf6), mRNA  NM_016604  Homo sapiens chromosome 5 open reading frame 5 (CSorf6), mRNA  NM_016605  Homo sapiens chromosome 1 open reading frame 21 (C11orf21), mRNA  NM_016605  Homo sapiens chromosome 1 open reading frame 21 (C11orf21), mRNA  NM_033508  Homo sapiens chromosome 1 open reading frame 21 (C11orf21), mRNA  NM_033507  Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 2, nuclear gene encoding mitochondrial protein, mRNA  NM_000162  Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 1, nuclear gene encoding mitochondrial protein, mRNA		
mitochondrial protein, mRNA  NM_000188 Homo sapiens hexokinase 1 (HK1), transcript variant 1, nuclear gene encoding mitochondrial protein, mRNA  NM_004728 Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 21 (DDX21), mRNA  NM_022148 Homo sapiens cytokine receptor-like factor 2 (CRLF2), mRNA  NM_022337 Homo sapiens RAB38, member RAS oncogene family (RAB38), mRNA  NM_016428 Homo sapiens NESH protein (NESH), mRNA  NM_016227 Homo sapiens chromosome 1 open reading frame 9 (Clorf9), mRNA  NM_014283 Homo sapiens chromosome 1 open reading frame 9 (Clorf9), mRNA  NM_014284 NM_014284 Homo sapiens TPA regulated locus (TPARL), mRNA  NM_030934 Homo sapiens chromosome 1 open reading frame 25 (Clorf25), mRNA  NM_030934 Homo sapiens chromosome 1 open reading frame 14 (Clorf14), mRNA  NM_030936 Homo sapiens chromosome 1 open reading frame 13 (Clorf13), mRNA  NM_016604 Homo sapiens chromosome 5 open reading frame 7 (C5orf7), mRNA  NM_016605 Homo sapiens chromosome 5 open reading frame 6 (C5orf6), mRNA  NM_016603 Homo sapiens chromosome 5 open reading frame 21 (C11orf21), mRNA  NM_014144 Homo sapiens chromosome 5 open reading frame 21 (C11orf21), mRNA  NM_033508 Homo sapiens chromosome 11 open reading frame 21 (C11orf21), mRNA  NM_033507 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 3, nuclear gene encoding mitochondrial protein, mRNA  NM_000162 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 2, nuclear gene encoding mitochondrial protein, mRNA  NM_000162 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 1, nuclear gene encoding mitochondrial protein, mRNA		<del> </del>
NM_000188 Homo sapiens hexokinase 1 (HK1), transcript variant 1, nuclear gene encoding mitochondrial protein, mRNA  NM_004728 Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 21 (DDX21), mRNA  NM_022148 Homo sapiens cytokine receptor-like factor 2 (CRLF2), mRNA  NM_022337 Homo sapiens RAB38, member RAS oncogene family (RAB38), mRNA  NM_016428 Homo sapiens NESH protein (NESH), mRNA  NM_016227 Homo sapiens chromosome 1 open reading frame 9 (C1orf9), mRNA  NM_014283 Homo sapiens chromosome 1 open reading frame 9 (C1orf9), mRNA  NM_018475 Homo sapiens TPA regulated locus (TPARL), mRNA  NM_020461 Homo sapiens gamma-tubulin complex component (GCP6), mRNA  NM_030934 Homo sapiens chromosome 1 open reading frame 25 (C1orf25), mRNA  NM_030935 Homo sapiens chromosome 1 open reading frame 13 (C1orf14), mRNA  NM_030936 Homo sapiens chromosome 1 open reading frame 13 (C1orf13), mRNA  NM_016604 Homo sapiens chromosome 5 open reading frame 6 (C5orf6), mRNA  NM_016605 Homo sapiens chromosome 5 open reading frame 6 (C5orf6), mRNA  NM_014144 Homo sapiens chromosome 5 open reading frame 2 (C1orf21), mRNA  NM_033508 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 3, nuclear gene encoding mitochondrial protein, mRNA  NM_000162 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 2, nuclear gene encoding mitochondrial protein, mRNA  NM_000162 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 2, nuclear gene encoding mitochondrial protein, mRNA		
mitochondrial protein, mRNA  NM_004728 Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 21 (DDX21), mRNA  NM_022148 Homo sapiens cytokine receptor-like factor 2 (CRLF2), mRNA  NM_022337 Homo sapiens RAB38, member RAS oncogene family (RAB38), mRNA  NM_016428 Homo sapiens NESH protein (NESH), mRNA  NM_016227 Homo sapiens chromosome 1 open reading frame 9 (C1orf9), mRNA  NM_014283 Homo sapiens chromosome 1 open reading frame 9 (C1orf9), mRNA  NM_018475 Homo sapiens TPA regulated locus (TPARL), mRNA  NM_020461 Homo sapiens gamma-tubulin complex component (GCP6), mRNA  NM_030934 Homo sapiens chromosome 1 open reading frame 25 (C1orf25), mRNA  NM_030933 Homo sapiens chromosome 1 open reading frame 13 (C1orf14), mRNA  NM_030936 Homo sapiens chromosome 1 open reading frame 13 (C1orf13), mRNA  NM_016604 Homo sapiens chromosome 5 open reading frame 7 (C5orf7), mRNA  NM_016605 Homo sapiens chromosome 5 open reading frame 6 (C5orf6), mRNA  NM_014144 Homo sapiens chromosome 1 open reading frame 21 (C11orf21), mRNA  NM_014144 Homo sapiens chromosome 5 open reading frame 21 (C11orf21), mRNA  NM_03508 Homo sapiens chromosome 1 open reading frame 21 (C11orf21), mRNA  NM_03508 Homo sapiens chromosome 5 open reading frame 21 (C11orf21), mRNA  NM_033508 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 3, nuclear gene encoding mitochondrial protein, mRNA  NM_000162 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 2, nuclear gene encoding mitochondrial protein, mRNA	NM 000188	
NM_022148 Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 21 (DDX21), mRNA  NM_022337 Homo sapiens cytokine receptor-like factor 2 (CRLF2), mRNA  NM_022337 Homo sapiens RAB38, member RAS oncogene family (RAB38), mRNA  NM_016428 Homo sapiens NESH protein (NESH), mRNA  NM_016227 Homo sapiens chromosome 1 open reading frame 9 (Clorf9), mRNA  NM_014283 Homo sapiens thromosome 1 open reading frame 9 (Clorf9), mRNA  NM_020461 Homo sapiens TPA regulated locus (TPARL), mRNA  NM_030934 Homo sapiens chromosome 1 open reading frame 25 (Clorf25), mRNA  NM_030933 Homo sapiens chromosome 1 open reading frame 14 (Clorf14), mRNA  NM_030934 Homo sapiens chromosome 1 open reading frame 13 (Clorf13), mRNA  NM_03069 Homo sapiens chromosome 1 open reading frame 13 (Clorf13), mRNA  NM_016604 Homo sapiens chromosome 5 open reading frame 6 (C5orf6), mRNA  NM_016605 Homo sapiens chromosome 5 open reading frame 6 (C5orf6), mRNA  NM_014144 Homo sapiens chromosome 1 open reading frame 5 (C5orf5), mRNA  NM_033508 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 2, nuclear gene encoding mitochondrial protein, mRNA  NM_000162 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 2, nuclear gene encoding mitochondrial protein, mRNA  NM_000162 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 2, nuclear gene encoding mitochondrial protein, mRNA		
NM_022148Homo sapiens cytokine receptor-like factor 2 (CRLF2), mRNANM_022337Homo sapiens RAB38, member RAS oncogene family (RAB38), mRNANM_016428Homo sapiens NESH protein (NESH), mRNANM_016227Homo sapiens chromosome 1 open reading frame 9 (C1orf9), mRNANM_014283Homo sapiens chromosome 1 open reading frame 9 (C1orf9), mRNANM_018475Homo sapiens TPA regulated locus (TPARL), mRNANM_020461Homo sapiens gamma-tubulin complex component (GCP6), mRNANM_030934Homo sapiens chromosome 1 open reading frame 25 (C1orf25), mRNANM_030933Homo sapiens chromosome 1 open reading frame 14 (C1orf14), mRNANM_030769Homo sapiens chromosome 5 open reading frame 13 (C1orf13), mRNANM_016604Homo sapiens chromosome 5 open reading frame 6 (C5orf5), mRNANM_016605Homo sapiens chromosome 5 open reading frame 6 (C5orf5), mRNANM_014144Homo sapiens chromosome 11 open reading frame 21 (C11orf21), mRNANM_033508Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 3, nuclear gene encoding mitochondrial protein, mRNANM_033507Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 2, nuclear gene encoding mitochondrial protein, mRNANM_000162Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 1, nuclear gene encoding mitochondrial protein, mRNA	NM_004728	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 21 (DDX21),
NM_022337Homo sapiens RAB38, member RAS oncogene family (RAB38), mRNANM_016428Homo sapiens NESH protein (NESH), mRNANM_016227Homo sapiens chromosome 1 open reading frame 9 (C1orf9), mRNANM_014283Homo sapiens chromosome 1 open reading frame 9 (C1orf9), mRNANM_018475Homo sapiens TPA regulated locus (TPARL), mRNANM_020461Homo sapiens gamma-tubulin complex component (GCP6), mRNANM_030934Homo sapiens chromosome 1 open reading frame 25 (C1orf25), mRNANM_030935Homo sapiens chromosome 1 open reading frame 13 (C1orf13), mRNANM_030769Homo sapiens chromosome 5 open reading frame 7 (C5orf7), mRNANM_016604Homo sapiens chromosome 5 open reading frame 6 (C5orf6), mRNANM_016605Homo sapiens chromosome 5 open reading frame 6 (C5orf6), mRNANM_016604Homo sapiens chromosome 1 open reading frame 21 (C11orf21), mRNANM_014144Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 3, nuclear gene encoding mitochondrial protein, mRNANM_033507Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 2, nuclear gene encoding mitochondrial protein, mRNANM_000162Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 1, nuclear gene encoding mitochondrial protein, mRNA	NM 022148	· · · · · · · · · · · · · · · · · · ·
NM_016428Homo sapiens NESH protein (NESH), mRNANM_016227Homo sapiens chromosome 1 open reading frame 9 (C1orf9), mRNANM_014283Homo sapiens chromosome 1 open reading frame 9 (C1orf9), mRNANM_018475Homo sapiens TPA regulated locus (TPARL), mRNANM_020461Homo sapiens gamma-tubulin complex component (GCP6), mRNANM_030934Homo sapiens chromosome 1 open reading frame 25 (C1orf25), mRNANM_030933Homo sapiens chromosome 1 open reading frame 14 (C1orf14), mRNANM_030769Homo sapiens chromosome 1 open reading frame 13 (C1orf13), mRNANM_016604Homo sapiens chromosome 5 open reading frame 6 (C5orf6), mRNANM_016605Homo sapiens chromosome 5 open reading frame 6 (C5orf6), mRNANM_014144Homo sapiens chromosome 11 open reading frame 21 (C11orf21), mRNANM_033508Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 3, nuclear gene encoding mitochondrial protein, mRNANM_033507Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 2, nuclear gene encoding mitochondrial protein, mRNANM_000162Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 1, nuclear gene encoding mitochondrial protein, mRNA	<del></del>	
NM_016227 Homo sapiens chromosome 1 open reading frame 9 (C1orf9), mRNA NM_014283 Homo sapiens chromosome 1 open reading frame 9 (C1orf9), mRNA NM_018475 Homo sapiens TPA regulated locus (TPARL), mRNA NM_020461 Homo sapiens gamma-tubulin complex component (GCP6), mRNA NM_030934 Homo sapiens chromosome 1 open reading frame 25 (C1orf25), mRNA NM_030933 Homo sapiens chromosome 1 open reading frame 14 (C1orf14), mRNA NM_030769 Homo sapiens chromosome 1 open reading frame 13 (C1orf13), mRNA NM_016604 Homo sapiens chromosome 5 open reading frame 7 (C5orf7), mRNA NM_016605 Homo sapiens chromosome 5 open reading frame 6 (C5orf6), mRNA NM_016603 Homo sapiens chromosome 5 open reading frame 5 (C5orf5), mRNA NM_014144 Homo sapiens chromosome 11 open reading frame 21 (C11orf21), mRNA NM_033508 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 3, nuclear gene encoding mitochondrial protein, mRNA  NM_033507 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 2, nuclear gene encoding mitochondrial protein, mRNA  NM_000162 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 1, nuclear gene encoding mitochondrial protein, mRNA		<del>                                      </del>
NM 014283 Homo sapiens chromosome 1 open reading frame 9 (C1orf9), mRNA  NM 018475 Homo sapiens TPA regulated locus (TPARL), mRNA  NM 020461 Homo sapiens gamma-tubulin complex component (GCP6), mRNA  NM 030934 Homo sapiens chromosome 1 open reading frame 25 (C1orf25), mRNA  NM 030933 Homo sapiens chromosome 1 open reading frame 14 (C1orf14), mRNA  NM 030769 Homo sapiens chromosome 1 open reading frame 13 (C1orf13), mRNA  NM 016604 Homo sapiens chromosome 5 open reading frame 7 (C5orf7), mRNA  NM 016605 Homo sapiens chromosome 5 open reading frame 6 (C5orf6), mRNA  NM 016603 Homo sapiens chromosome 5 open reading frame 5 (C5orf5), mRNA  NM 014144 Homo sapiens chromosome 11 open reading frame 21 (C11orf21), mRNA  NM 033508 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 3, nuclear gene encoding mitochondrial protein, mRNA  NM 000162 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 2, nuclear gene encoding mitochondrial protein, mRNA  NM 000162 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 1, nuclear gene encoding mitochondrial protein, mRNA		
NM_018475Homo sapiens TPA regulated locus (TPARL), mRNANM_020461Homo sapiens gamma-tubulin complex component (GCP6), mRNANM_030934Homo sapiens chromosome 1 open reading frame 25 (C1orf25), mRNANM_030933Homo sapiens chromosome 1 open reading frame 14 (C1orf14), mRNANM_030769Homo sapiens chromosome 1 open reading frame 13 (C1orf13), mRNANM_016604Homo sapiens chromosome 5 open reading frame 7 (C5orf7), mRNANM_016605Homo sapiens chromosome 5 open reading frame 6 (C5orf6), mRNANM_016603Homo sapiens chromosome 1 open reading frame 5 (C5orf5), mRNANM_014144Homo sapiens chromosome 11 open reading frame 21 (C11orf21), mRNANM_033508Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 3, nuclear gene encoding mitochondrial protein, mRNANM_033507Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 2, nuclear gene encoding mitochondrial protein, mRNANM_000162Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 1, nuclear gene encoding mitochondrial protein, mRNA		
NM 020461 Homo sapiens gamma-tubulin complex component (GCP6), mRNA NM 030934 Homo sapiens chromosome 1 open reading frame 25 (C1orf25), mRNA NM 030933 Homo sapiens chromosome 1 open reading frame 14 (C1orf14), mRNA NM 030769 Homo sapiens chromosome 1 open reading frame 13 (C1orf13), mRNA NM 016604 Homo sapiens chromosome 5 open reading frame 7 (C5orf7), mRNA NM 016605 Homo sapiens chromosome 5 open reading frame 6 (C5orf6), mRNA NM 016603 Homo sapiens chromosome 5 open reading frame 5 (C5orf5), mRNA NM 014144 Homo sapiens chromosome 11 open reading frame 21 (C11orf21), mRNA NM 033508 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 3, nuclear gene encoding mitochondrial protein, mRNA  NM 000162 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 2, nuclear gene encoding mitochondrial protein, mRNA  NM 000162 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 1, nuclear gene encoding mitochondrial protein, mRNA		
NM 030934 Homo sapiens chromosome 1 open reading frame 25 (C1orf25), mRNA NM 030933 Homo sapiens chromosome 1 open reading frame 14 (C1orf14), mRNA NM 030769 Homo sapiens chromosome 1 open reading frame 13 (C1orf13), mRNA NM 016604 Homo sapiens chromosome 5 open reading frame 7 (C5orf7), mRNA NM 016605 Homo sapiens chromosome 5 open reading frame 6 (C5orf6), mRNA NM 016603 Homo sapiens chromosome 5 open reading frame 5 (C5orf5), mRNA NM 014144 Homo sapiens chromosome 11 open reading frame 21 (C11orf21), mRNA NM 033508 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 3, nuclear gene encoding mitochondrial protein, mRNA  NM 000162 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 2, nuclear gene encoding mitochondrial protein, mRNA  NM 000162 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 1, nuclear gene encoding mitochondrial protein, mRNA		<del> </del>
NM 030933 Homo sapiens chromosome 1 open reading frame 14 (C1orf14), mRNA NM 030769 Homo sapiens chromosome 1 open reading frame 13 (C1orf13), mRNA NM 016604 Homo sapiens chromosome 5 open reading frame 7 (C5orf7), mRNA NM 016605 Homo sapiens chromosome 5 open reading frame 6 (C5orf6), mRNA NM 016603 Homo sapiens chromosome 5 open reading frame 5 (C5orf5), mRNA NM 014144 Homo sapiens chromosome 11 open reading frame 21 (C11orf21), mRNA NM 033508 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 3, nuclear gene encoding mitochondrial protein, mRNA  NM 033507 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 2, nuclear gene encoding mitochondrial protein, mRNA  NM 000162 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 1, nuclear gene encoding mitochondrial protein, mRNA		
NM_030769 Homo sapiens chromosome 1 open reading frame 13 (C1orf13), mRNA  NM_016604 Homo sapiens chromosome 5 open reading frame 7 (C5orf7), mRNA  NM_016605 Homo sapiens chromosome 5 open reading frame 6 (C5orf6), mRNA  NM_016603 Homo sapiens chromosome 5 open reading frame 5 (C5orf5), mRNA  NM_014144 Homo sapiens chromosome 11 open reading frame 21 (C11orf21), mRNA  NM_033508 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 3, nuclear gene encoding mitochondrial protein, mRNA  NM_033507 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 2, nuclear gene encoding mitochondrial protein, mRNA  NM_000162 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 1, nuclear gene encoding mitochondrial protein, mRNA		
NM_016604 Homo sapiens chromosome 5 open reading frame 7 (C5orf7), mRNA  NM_016605 Homo sapiens chromosome 5 open reading frame 6 (C5orf6), mRNA  NM_016603 Homo sapiens chromosome 5 open reading frame 5 (C5orf5), mRNA  NM_014144 Homo sapiens chromosome 11 open reading frame 21 (C11orf21), mRNA  NM_033508 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 3, nuclear gene encoding mitochondrial protein, mRNA  NM_033507 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 2, nuclear gene encoding mitochondrial protein, mRNA  NM_000162 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 1, nuclear gene encoding mitochondrial protein, mRNA		
NM_016605 Homo sapiens chromosome 5 open reading frame 6 (C5orf6), mRNA  NM_016603 Homo sapiens chromosome 5 open reading frame 5 (C5orf5), mRNA  NM_014144 Homo sapiens chromosome 11 open reading frame 21 (C11orf21), mRNA  NM_033508 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 3, nuclear gene encoding mitochondrial protein, mRNA  NM_033507 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 2, nuclear gene encoding mitochondrial protein, mRNA  NM_000162 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 1, nuclear gene encoding mitochondrial protein, mRNA		
NM_016603 Homo sapiens chromosome 5 open reading frame 5 (C5orf5), mRNA  NM_014144 Homo sapiens chromosome 11 open reading frame 21 (C11orf21), mRNA  NM_033508 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 3, nuclear gene encoding mitochondrial protein, mRNA  NM_033507 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 2, nuclear gene encoding mitochondrial protein, mRNA  NM_000162 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 1, nuclear gene encoding mitochondrial protein, mRNA		
NM_014144 Homo sapiens chromosome 11 open reading frame 21 (C11orf21), mRNA  NM_033508 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 3, nuclear gene encoding mitochondrial protein, mRNA  NM_033507 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 2, nuclear gene encoding mitochondrial protein, mRNA  NM_000162 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 1, nuclear gene encoding mitochondrial protein, mRNA		
NM_033508 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 3, nuclear gene encoding mitochondrial protein, mRNA  NM_033507 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 2, nuclear gene encoding mitochondrial protein, mRNA  NM_000162 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 1, nuclear gene encoding mitochondrial protein, mRNA		Homo sapiens chromosome 5 open reading frame 5 (C5orf5), mRNA
2) (GCK), transcript variant 3, nuclear gene encoding mitochondrial protein, mRNA  NM_033507  Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 2, nuclear gene encoding mitochondrial protein, mRNA  NM_000162  Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 1, nuclear gene encoding mitochondrial protein, mRNA		
2) (GCK), transcript variant 2, nuclear gene encoding mitochondrial protein, mRNA  NM_000162 Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young 2) (GCK), transcript variant 1, nuclear gene encoding mitochondrial protein, mRNA	NM_033508	2) (GCK), transcript variant 3, nuclear gene encoding mitochondrial protein,
2) (GCK), transcript variant 1, nuclear gene encoding mitochondrial protein, mRNA	NM_033507	2) (GCK), transcript variant 2, nuclear gene encoding mitochondrial protein,
	_	2) (GCK), transcript variant 1, nuclear gene encoding mitochondrial protein, mRNA
NM_025241 Homo sapiens UBX domain-containing 1 (UBXD1), mRNA		Homo sapiens UBX domain-containing 1 (UBXD1), mRNA
NM_002098 Homo sapiens guanylate cyclase activator 1B (retina) (GUCA1B), mRNA		
NM_003137 Homo sapiens SFRS protein kinase 1 (SRPK1), mRNA	NM_003137	Homo sapiens SFRS protein kinase 1 (SRPK1), mRNA
NM_003064 Homo sapiens secretory leukocyte protease inhibitor (antileukoproteinase) (SLPI), mRNA	NM_003064	
NM_033484 Homo sapiens F-box only protein 4 (FBXO4), transcript variant 2, mRNA	NM_033484	<del>                                      </del>

NM_012176	Homo sapiens F-box only protein 4 (FBXO4), transcript variant 1, mRNA
NM_000400	Homo sapiens excision repair cross-complementing rodent repair deficiency,
<del>-</del>	complementation group 2 (xeroderma pigmentosum D) (ERCC2), mRNA
NM 014266	Homo sapiens DNAX-activation protein 10 (DAP10), mRNA
NM 002821	Homo sapiens PTK7 protein tyrosine kinase 7 (PTK7), mRNA
NM 033502	Homo sapiens transcriptional regulating protein 132 (TReP-132), transcript
_	variant 1, mRNA
NM_033501	Homo sapiens transcriptional regulating protein 132 (TReP-132), transcript
	variant 2, mRNA
NM_018415	Homo sapiens transcriptional regulating protein 132 (TReP-132), transcript
	variant 3, mRNA
NM_000994	Homo sapiens ribosomal protein L32 (RPL32), mRNA
NM_033437	Homo sapiens phosphodiesterase 5A, cGMP-specific (PDE5A), transcript variant
	3, mRNA
NM_033431	Homo sapiens phosphodiesterase 5A, cGMP-specific (PDE5A), transcript variant
	4, mRNA
NM_033430	Homo sapiens phosphodiesterase 5A, cGMP-specific (PDE5A), transcript variant
	2, mRNA
NM_001083	Homo sapiens phosphodiesterase 5A, cGMP-specific (PDE5A), transcript variant
	1, mRNA
NM_000189	Homo sapiens hexokinase 2 (HK2), mRNA
NM_033185	Homo sapiens keratin associated protein 3.3 (KAP3.3), mRNA
NM_031959	Homo sapiens keratin associated protein 3.2 (KRTAP3.2), mRNA
NM_033481	Homo sapiens F-box only protein 9 (FBXO9), transcript variant 3, mRNA
NM_033480	Homo sapiens F-box only protein 9 (FBXO9), transcript variant 2, mRNA
NM_012347	Homo sapiens F-box only protein 9 (FBXO9), transcript variant 1, mRNA
NM_033506	Homo sapiens F-box only protein 24 (FBXO24), transcript variant 1, mRNA Homo sapiens F-box only protein 24 (FBXO24), transcript variant 2, mRNA
NM_012172	Homo sapiens F-box only protein 7 (FBXO24), transcript variant 2, income Homo sapiens F-box only protein 7 (FBXO7), mRNA
NM_012179	Homo sapiens F-box only protein 7 (FBXO7), mRNA  Homo sapiens F-box only protein 6 (FBXO6), mRNA
NM_018438	Homo sapiens F-box only protein 5 (FBXO5), mRNA
NM_012177	Homo sapiens F-box only protein 3 (FBXO30), mRNA
NM_032145	Homo sapiens r-box protein 30 (PBAO30), interval  Homo sapiens a disintegrin and metalloproteinase domain 21 (ADAM21),
NM_003813	mRNA
NM_003814	Homo sapiens a disintegrin and metalloproteinase domain 20 (ADAM20),
11111_003614	mRNA
NM 015698	Homo sapiens T54 protein (T54), mRNA
NM 033222	Homo sapiens PC4 and SFRS1 interacting protein 2 (PSIP2), mRNA
NM 002887	Homo sapiens arginyl-tRNA synthetase (RARS), mRNA
NM 033084	Homo sapiens Fanconi anemia, complementation group D2 (FANCD2), mRNA
NM 014005	Homo sapiens protocadherin alpha 9 (PCDHA9), transcript variant 2, mRNA
NM 018902	Homo sapiens protocadherin alpha 11 (PCDHA11), transcript variant 1, mRNA
NM 031882	Homo sapiens protocadherin alpha subfamily C, 1 (PCDHAC1), transcript
1111_051002	variant 2, mRNA
NM 018898	Homo sapiens protocadherin alpha subfamily C, 1 (PCDHAC1), transcript
1111_010070	variant 1, mRNA
NM 031883	Homo sapiens protocadherin alpha subfamily C, 2 (PCDHAC2), transcript
1111_051005	variant 2, mRNA
NM 018899	Homo sapiens protocadherin alpha subfamily C, 2 (PCDHAC2), transcript
	variant 1, mRNA
NM 019119	Homo sapiens protocadherin beta 9 (PCDHB9), mRNA
NM 018916	Homo sapiens protocadherin gamma subfamily A, 3 (PCDHGA3), transcript
	1

	variant 1, mRNA
NM 032704	Homo sapiens tubulin alpha 6 (TUBA6), mRNA
NM_032407	Homo sapiens protocadherin gamma subfamily C, 5 (PCDHGC5), transcript
1111_032.07	variant 2, mRNA
NM_018929	Homo sapiens protocadherin gamma subfamily C, 5 (PCDHGC5), transcript
	variant 1, mRNA
NM_032406	Homo sapiens protocadherin gamma subfamily C, 4 (PCDHGC4), transcript
<u>-</u>	variant 2, mRNA
NM_018928	Homo sapiens protocadherin gamma subfamily C, 4 (PCDHGC4), transcript
_	variant 1, mRNA
NM 032101	Homo sapiens protocadherin gamma subfamily B, 7 (PCDHGB7), transcript
_	variant 2, mRNA
NM_018927	Homo sapiens protocadherin gamma subfamily B, 7 (PCDHGB7), transcript
<del>_</del>	variant 1, mRNA
NM_032099	Homo sapiens protocadherin gamma subfamily B, 5 (PCDHGB5), transcript
	variant 2, mRNA
NM_018925	Homo sapiens protocadherin gamma subfamily B, 5 (PCDHGB5), transcript
-	variant 1, mRNA
NM_032100	Homo sapiens protocadherin gamma subfamily B, 6 (PCDHGB6), transcript
	variant 2, mRNA
NM_018926	Homo sapiens protocadherin gamma subfamily B, 6 (PCDHGB6), transcript
>P 5 00000	variant 1, mRNA
NM_032097	Homo sapiens protocadherin gamma subfamily B, 3 (PCDHGB3), transcript
ND ( 019024	variant 2, mRNA
NM_018924	Homo sapiens protocadherin gamma subfamily B, 3 (PCDHGB3), transcript variant 1, mRNA
NM 032096	Homo sapiens protocadherin gamma subfamily B, 2 (PCDHGB2), transcript
14WI_032030	variant 2, mRNA
NM_018923	Homo sapiens protocadherin gamma subfamily B, 2 (PCDHGB2), transcript
	variant 1, mRNA
NM_032095	Homo sapiens protocadherin gamma subfamily B, 1 (PCDHGB1), transcript
	variant 2, mRNA
NM_018922	Homo sapiens protocadherin gamma subfamily B, 1 (PCDHGB1), transcript
	variant 1, mRNA
NM_032089	Homo sapiens protocadherin gamma subfamily A, 9 (PCDHGA9), transcript
	variant 2, mRNA
NM_018921	Homo sapiens protocadherin gamma subfamily A, 9 (PCDHGA9), transcript
	variant 1, mRNA
NM_032088	Homo sapiens protocadherin gamma subfamily A, 8 (PCDHGA8), transcript
	variant 1, mRNA
NM_014004	Homo sapiens protocadherin gamma subfamily A, 8 (PCDHGA8), transcript
ND 6 000055	variant 2, mRNA
NM_032853	Homo sapiens hypothetical protein FLJ14868 (FLJ14868), mRNA
NM_032589	Homo sapiens Down syndrome critical region gene 8 (DSCR8), mRNA
NM_032087	
NIM 018020	
14141 01 0250	
NIM 032086	
14141_025000	
NM 018919	
11112_010717	
NM_032087 NM_018920 NM_032086 NM_018919	Homo sapiens protocadherin gamma subfamily A, 7 (PCDHGA7), transcript variant 2, mRNA  Homo sapiens protocadherin gamma subfamily A, 7 (PCDHGA7), transcript variant 1, mRNA  Homo sapiens protocadherin gamma subfamily A, 6 (PCDHGA6), transcript variant 2, mRNA  Homo sapiens protocadherin gamma subfamily A, 6 (PCDHGA6), transcript variant 2, mRNA  Homo sapiens protocadherin gamma subfamily A, 6 (PCDHGA6), transcript variant 1, mRNA

cript cript cript cript cript cript cript
eript eript eript
eript
cript
cript
cript
cript
nscript
cript
anscript
anscript
anscript
anscript
DNA
nRNA
nRNA nRNA

NM_031497	Homo sapiens protocadherin alpha 3 (PCDHA3), transcript variant 2, mRNA
NM_018906	Homo sapiens protocadherin alpha 3 (PCDHA3), transcript variant 1, mRNA
NM_031496	Homo sapiens protocadherin alpha 2 (PCDHA2), transcript variant 3, mRNA
NM_031495	Homo sapiens protocadherin alpha 2 (PCDHA2), transcript variant 2, mRNA
NM_018905	Homo sapiens protocadherin alpha 2 (PCDHA2), transcript variant 1, mRNA
NM_031411	Homo sapiens protocadherin alpha 1 (PCDHA1), transcript variant 3, mRNA
NM_031410	Homo sapiens protocadherin alpha 1 (PCDHA1), transcript variant 2, mRNA
NM_018900	Homo sapiens protocadherin alpha 1 (PCDHA1), transcript variant 1, mRNA
NM_031865	Homo sapiens protocadherin alpha 13 (PCDHA13), transcript variant 2, mRNA
NM_018904	Homo sapiens protocadherin alpha 13 (PCDHA13), transcript variant 1, mRNA
NM_031849	Homo sapiens protocadherin alpha 6 (PCDHA6), transcript variant 3, mRNA
NM 031864	Homo sapiens protocadherin alpha 12 (PCDHA12), transcript variant 2, mRNA
NM 031848	Homo sapiens protocadherin alpha 6 (PCDHA6), transcript variant 2, mRNA
NM 018903	Homo sapiens protocadherin alpha 12 (PCDHA12), transcript variant 1, mRNA
NM 031861	Homo sapiens protocadherin alpha 11 (PCDHA11), transcript variant 2, mRNA
NM 018909	Homo sapiens protocadherin alpha 6 (PCDHA6), transcript variant 1, mRNA
NM 031860	Homo sapiens protocadherin alpha 10 (PCDHA10), transcript variant 3, mRNA
NM 031859	Homo sapiens protocadherin alpha 10 (PCDHA10), transcript variant 2, mRNA
NM 018901	Homo sapiens protocadherin alpha 10 (PCDHA10), transcript variant 1, mRNA
NM 015429	Homo sapiens DKFZP586L2024 protein (NESHBP), mRNA
NM 031481	Homo sapiens solute carrier family 25, (mitochondrial carrier), member 18
	(SLC25A18), mRNA
NM 031442	Homo sapiens brain cell membrane protein 1 (BCMP1), mRNA
NM_030762	Homo sapiens basic helix-loop-helix domain containing, class B, 3 (BHLHB3),
	mRNA
NM 023035	Homo sapiens calcium channel, voltage-dependent, P/Q type, alpha 1A subunit
	(CACNA1A), transcript variant 2, mRNA
NM 014487	Homo sapiens nucleolar cysteine-rich protein (HSA6591), mRNA
NM 025239	Homo sapiens programmed death ligand 2 (PDL2), mRNA
NM 024859	Homo sapiens hypothetical protein FLJ21687 (FLJ21687), mRNA
NM 000575	Homo sapiens interleukin 1, alpha (IL1A), mRNA
NM 005348	Homo sapiens heat shock 90kD protein 1, alpha (HSPCA), mRNA
NM 006900	Homo sapiens interferon, alpha 13 (IFNA13), mRNA
NM 023067	Homo sapiens forkhead transcription factor FOXL2 (BPES), mRNA
NM 022552	Homo sapiens DNA (cytosine-5-)-methyltransferase 3 alpha (DNMT3A), mRNA
NM 022346	Homo sapiens chromosome condensation protein G (HCAP-G), mRNA
NM 022119	Homo sapiens protease, serine, 22 (PRSS22), mRNA
NM 022062	Homo sapiens PBX/knotted 1 homeobox 2 (PKNOX2), mRNA
NM 018665	Homo sapiens DEAD-box protein (HAGE), mRNA
NM 004614	Homo sapiens thymidine kinase 2, mitochondrial (TK2), mRNA
NM 020346	Homo sapiens solute carrier family 17 (sodium-dependent inorganic phosphate
	cotransporter), member 6 (SLC17A6), mRNA
NM_020309	Homo sapiens solute carrier family 17 (sodium-dependent inorganic phosphate
2.1.1_020309	cotransporter), member 7 (SLC17A7), mRNA
NM 020131	Homo sapiens chromosome 1 open reading frame 6 (Clorf6), mRNA
NM 017444	Homo sapiens chromatin accessibility complex 1 (CHRAC1), mRNA
NM 016260	Homo sapiens zinc finger protein, subfamily 1A, 2 (Helios) (ZNFN1A2), mRNA
NM 015510	Homo sapiens DKFZP566O084 protein (DKFZp566O084), mRNA
NM 014433	Homo sapiens rhabdoid tumor deletion region gene 1 (RTDR1), mRNA
NM 014312	Homo sapiens cortical thymocyte receptor (X. laevis CTX) like (CTXL), mRNA
NM 004539	Homo sapiens asparaginyl-tRNA synthetase (NARS), mRNA
NM 013284	Homo sapiens asparagmyr-txiva synthetase (tvaxes), mixtxi  Homo sapiens polymerase (DNA directed), mu (POLM), mRNA
14141_013264	I nomo sapiens porymerase (Diva directed), ma (1 ODN), ma vi

	The state of the s
NM_013274	Homo sapiens polymerase (DNA directed), lambda (POLL), mRNA
NM_003235	Theme conjugate thyroglobulin (1(t), mKNA
NM_001963	Homo sapiens epidermal growth factor (beta-urogastrone) (EGF), mRNA
NM_007158	Homo sapiens NRAS-related gene (D1S155E), mRNA
NM_007000	Homo sapiens uroplakin 1A (UPK1A), mRNA
NM_006947	Homo sapiens signal recognition particle 72kD (SRP72), mRNA  Homo sapiens signal recognition particle 72kD (SRP72), mRNA
NM_006892	Homo sapiens DNA (cytosine-5-)-methyltransferase 3 beta (DNMT3B), mRNA
NM_006760	Homo sapiens uroplakin 2 (UPK2), mRNA
NM_006691	Homo sapiens extracellular link domain-containing 1 (XLKD1), mRNA  Homo sapiens extracellular link domain-containing 1 (XLKD1), mRNA
NM_006572	Homo sapiens guanine nucleotide binding protein (G protein), alpha 13 (GNA13), mRNA
NM 006494	Homo saniens Ets2 repressor factor (ERF), mRNA
NM_006352	Homo saniens zinc finger protein 238 (ZNF238), mRNA
NM 006082	Homo capiens tubulin alpha ubiquitous (K-ALPHA-1), mknA
NM_005084	Homo sapiens phospholipase A2, group VII (platelet-activating factor
14141_00500:	acetylhydrolase, plasma) (PLA2G7), mRNA
NM 004999	Homo saniens myosin VI (MYO6), mRNA
NM 004937	Home genions systingsis nephronathic (CTNS), mRNA
NM_004212	Homo sapiens solute carrier family 28 (sodium-coupled nucleoside transporter),
ND4 004555	Homo sapiens nuclear factor of activated T-cells, cytoplasmic, calcineurin-
NM_004555	dependent 3 (NEATC3) mRNA
NM_004554	Homo sapiens nuclear factor of activated T-cells, cytoplasmic, calcineurin-
NWI_004334	January don't 4 (NTEATCA) mRNA
NM_000695	Homo sapiens aldehyde dehydrogenase 3 family, member B2 (ALDH3B2),
14141_000075	
NM_000373	Homo saniens uridine monophosphate synthetase (orotate phosphoribosyl
14141_000375	transferace and protidine-5'-decarboxylase) (UMPS), mknA
NM_003332	Homo sapiens TYRO protein tyrosine kinase binding protein (TYROBP),
11111	mRNA
NM 000367	Homo sapiens thiopurine S-methyltransferase (TPMT), mRNA
NM_001250	Homo sapiens tumor necrosis factor receptor superfamily, member 5 (TNFRSF5), mRNA
ND4 002990	Homo sapiens v-raf-1 murine leukemia viral oncogene homolog 1 (RAF1),
NM_002880	DNIA
NM_003978	Homo sapiens proline-serine-threonine phosphatase interacting protein I
	(PSTPIP1), mRNA  Homo sapiens prostate cancer overexpressed gene 1 (POV1), mRNA
NM_003627	Homo sapiens oviductal glycoprotein 1, 120kD (mucin 9, oviductin) (OVGP1),
NM_002557	
77.5.000541	mRNA Homo sapiens oxoglutarate (alpha-ketoglutarate) dehydrogenase (lipoamide)
NM_002541	(OGDH), mRNA
377 6 000 40 6	Homo sapiens gonadotropin-releasing hormone receptor (GNRHR), mRNA
NM_000406 NM_001979	Homo sapiens epoxide hydrolase 2, cytoplasmic (EPHX2), mRNA
	Homo seniens cyclin F (CCNF), mRNA
NM_001761 NM_001190	
NM_001130	mPNA
NM_000485	1 1 C (ADDT) DNA
NM 033514	
NM_033495	TImes serious VIA A 1309 protein (KIAAI309), mRNA
NM_022436	
INIVI_022430	1) (ABCG5), mRNA
	1 -7 (-1 - 0 -7), =

NM_016333	Homo sapiens serine/arginine repetitive matrix 2 (SRRM2), mRNA
NM_012412	Homo sapiens histone H2A.F/Z variant (H2AV), mRNA
NM_001897	Homo sapiens chondroitin sulfate proteoglycan 4 (melanoma-associated)
	(CSPG4), mRNA
NM_031420	Homo sapiens mitochondrial ribosomal protein L9 (MRPL9), mRNA
NM_020393	Homo sapiens hypothetical protein SBBI67 (LOC57115), mRNA
NM_015956	Homo sapiens mitochondrial ribosomal protein L4 (MRPL4), mRNA
NM_004537	Homo sapiens nucleosome assembly protein 1-like 1 (NAP1L1), mRNA
NM_033504	Homo sapiens CAC-1 (CAC-1), mRNA
NM_033503	Homo sapiens Bcl-2 modifying factor (BMF), mRNA
NM_022059	Homo sapiens chemokine (C-X-C motif) ligand 16 (CXCL16), mRNA
NM_022048	Homo sapiens casein kinase 1, gamma 1 (CSNK1G1), mRNA
NM_019009	Homo sapiens Toll-interacting protein (TOLLIP), mRNA
NM_018058	Homo sapiens cartilage acidic protein 1 (CRTAC1), mRNA
NM_017443	Homo sapiens polymerase (DNA directed), epsilon 3 (p17 subunit) (POLE3), mRNA
NM_007359	Homo sapiens MLN51 protein (MLN51), mRNA
NM 030956	Homo sapiens toll-like receptor 10 (TLR10), mRNA
NM 020653	Homo sapiens zinc finger protein 287 (ZNF287), mRNA
NM 020652	Homo sapiens zinc finger protein 286 (ZNF286), mRNA
NM 020365	Homo sapiens eukaryotic translation initiation factor 2B, subunit 3 (gamma,
	58kD) (EIF2B3), mRNA
NM_013432	Homo sapiens nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor-like 2 (NFKBIL2), mRNA
NM_003740	Homo sapiens potassium channel, subfamily K, member 5 (TASK-2) (KCNK5), mRNA
NM_033311	Homo sapiens potassium inwardly-rectifying channel, subfamily K, member 4 (KCNK4), transcript variant 3, mRNA
NM_033310	Homo sapiens potassium inwardly-rectifying channel, subfamily K, member 4 (KCNK4), transcript variant 2, mRNA
NM_016611	Homo sapiens potassium inwardly-rectifying channel, subfamily K, member 4 (KCNK4), transcript variant 1, mRNA
NM_033360	Homo sapiens v-Ki-ras2 Kirsten rat sarcoma 2 viral oncogene homolog (KRAS2), transcript variant a, mRNA
NM_004985	Homo sapiens v-Ki-ras2 Kirsten rat sarcoma 2 viral oncogene homolog (KRAS2), transcript variant b, mRNA
NM_022442	Homo sapiens ubiquitin-conjugating enzyme E2 variant 1 (UBE2V1), transcript variant 3, mRNA
NM_021988	Homo sapiens ubiquitin-conjugating enzyme E2 variant 1 (UBE2V1), transcript variant 1, mRNA
NM_003349	Homo sapiens ubiquitin-conjugating enzyme E2 variant 1 (UBE2V1), transcript variant 2, mRNA
NM 003546	Homo sapiens H4 histone family, member K (H4FK), mRNA
NM 003541	Homo sapiens H4 histone family, member D (H4FD), mRNA
NM_003536	Homo sapiens H3 histone family, member K (H3FK), mRNA
NM_003535	Homo sapiens H3 histone family, member J (H3FJ), mRNA
NM_003533	Homo sapiens H3 histone family, member F (H3FF), mRNA
NM_003521	Homo sapiens H2B histone family, member E (H2BFE), mRNA
NM_003520	Homo sapiens H2B histone family, member D (H2BFD), mRNA
NM_003519	Homo sapiens H2B histone family, member C (H2BFC), mRNA
NM 003514	Homo sapiens H2A histone family, member N (H2AFN), mRNA
NM_003511	Homo sapiens H2A histone family, member I (H2AFI), mRNA
:	1

······································	C (HIPS) DAIA
NM_005322	Homo sapiens H1 histone family, member 5 (H1F5), mRNA
NM_021066	Homo sapiens H2A histone family, member E (H2AFE), mRNA
NM_003510	Homo sapiens H2A histone family, member D (H2AFD), mRNA
NM_003509	Homo sapiens H2A histone family, member C (H2AFC), mRNA
NM_033358	Homo sapiens caspase 8, apoptosis-related cysteine protease (CASP8), transcript
	variant E, mRNA
NM_033357	Homo sapiens caspase 8, apoptosis-related cysteine protease (CASP8), transcript variant D, mRNA
NM_033356	Homo sapiens caspase 8, apoptosis-related cysteine protease (CASP8), transcript
	variant C, mRNA  Homo sapiens caspase 8, apoptosis-related cysteine protease (CASP8), transcript
NM_033355	variant B, mRNA
NM_001228	Homo sapiens caspase 8, apoptosis-related cysteine protease (CASP8), transcript
14141_001228	variant A, mRNA
ND 4 022240	Homo sapiens caspase 7, apoptosis-related cysteine protease (CASP7), transcript
NM_033340	variant beta mRNA
NM_033339	Homo sapiens caspase 7, apoptosis-related cysteine protease (CASP7), transcript
14141-02222	variant camma mRNA
NIM 022228	Homo sapiens caspase 7, apoptosis-related cysteine protease (CASP7), transcript
NM_033338	variant delta, mRNA
NM 001227	Homo sapiens caspase 7, apoptosis-related cysteine protease (CASP7), transcript
14141_001227	variant alpha, mRNA
NM 001005	Homo sapiens ribosomal protein S3 (RPS3), mRNA
NM 006013	Homo sapiens ribosomal protein L10 (RPL10), mRNA
NM 013368	Homo sapiens RPA-binding trans-activator (RBT1), mRNA
	Homo sapiens lymphocyte-activation gene 3 (LAG3), mRNA
NM_002286	Homo sapiens IL2-inducible T-cell kinase (ITK), mRNA
NM_005546	Homo sapiens it.2-inductore 1-cen kmase (11K), mktv1
NM_005538	Homo sapiens inhibin, beta C (INHBC), mRNA
NM_033257	Homo sapiens DiGeorge syndrome critical region gene 6 like (DGCR6L), mRNA
NM 001917	Homo sapiens D-amino-acid oxidase (DAO), mRNA
NM_001629	Homo sapiens arachidonate 5-lipoxygenase-activating protein (ALOX5AP), mRNA
NM 000024	Homo sapiens adrenergic, beta-2-, receptor, surface (ADRB2), mRNA
	Homo sapiens adrenergic, alpha-2C-, receptor (ADRA2C), mRNA
NM_000683	Homo sapiens adrenergic, alpha-28-, receptor (ADRA2B), mRNA
NM_000682	Homo sapiens adrenergic, alpha-2A-, receptor (ADRA2A), mRNA
NM_000681	Homo sapiens autenergie, aipita-2A-, receptor (ADIVEZI), find the
NM_006179	Homo sapiens neurotrophin 5 (neurotrophin 4/5) (NTF5), mRNA
NM_033277	Homo sapiens lacritin (LACRT), mRNA
NM_022128	Homo sapiens ribokinase (RBSK), mRNA
NM_004823	Homo sapiens potassium channel, subfamily K, member 6 (TWIK-2) (KCNK6), mRNA
NM 002246	Homo sapiens potassium channel, subfamily K, member 3 (TASK-1) (KCNK3),
1111_002210	mRNA
NM_032405	Homo sapiens transmembrane protease, serine 3 (TMPRSS3), transcript variant
14141_032403	D. mRNA
NM_032404	Homo sapiens transmembrane protease, serine 3 (TMPRSS3), transcript variant
	C, mRNA
NM_032401	Homo sapiens transmembrane protease, serine 3 (TMPRSS3), transcript variant B, mRNA
NM 024022	Homo sapiens transmembrane protease, serine 3 (TMPRSS3), transcript variant
	A, mRNA

NM 016234	Homo sapiens fatty-acid-Coenzyme A ligase, long-chain 5 (FACL5), mRNA
NM 006883	Homo sapiens short stature homeobox (SHOX), transcript variant SHOXb,
_	mRNA
NM_000451	Homo sapiens short stature homeobox (SHOX), transcript variant SHOXa,
	mRNA
NM_006476	Homo sapiens ATP synthase, H+ transporting, mitochondrial F0 complex,
	subunit g (ATP5L), mRNA
NM_006356	Homo sapiens ATP synthase, H+ transporting, mitochondrial F0 complex,
_	subunit d (ATP5H), mRNA
NM 024683	Homo sapiens hypothetical protein FLJ22729 (FLJ22729), mRNA
NM 033468	Homo sapiens zinc finger protein 257 (ZNF257), mRNA
NM 033453	Homo sapiens inosine triphosphatase (nucleoside triphosphate pyrophosphatase)
_	(ITPA), mRNA
NM 032144	Homo sapiens RAB6C, member RAS oncogene family (RAB6C), mRNA
NM 031296	Homo sapiens RAB33B, member RAS oncogene family (RAB33B), mRNA
NM 022570	Homo sapiens C-type (calcium dependent, carbohydrate-recognition domain)
1111_0000	lectin, superfamily member 12 (CLECSF12), mRNA
NM 022825	Homo sapiens porcupine (MG61), mRNA
NM 022449	Homo sapiens RAB17, member RAS oncogene family (RAB17), mRNA
NM 016322	Homo sapiens RAB14, member RAS oncogene family (RAB14), mRNA
NM 006331	Homo sapiens C2f protein (C2F), mRNA
NM 007066	Homo sapiens protein kinase (cAMP-dependent, catalytic) inhibitor gamma
1411_007000	(PKIG), mRNA
NM 002732	Homo sapiens protein kinase, cAMP-dependent, catalytic, gamma (PRKACG),
NWI_002732	mRNA
NM_005055	Homo sapiens receptor-associated protein of the synapse, 43kD (RAPSN),
14M_002022	transcript variant 1, mRNA
NM_032645	Homo sapiens receptor-associated protein of the synapse, 43kD (RAPSN),
14141_052045	transcript variant 2, mRNA
NM 033305	Homo sapiens chorea acanthocytosis (CHAC), transcript variant A, mRNA
NM 015186	Homo sapiens chorea acanthocytosis (CHAC), transcript variant B, mRNA
NM 004624	Homo sapiens vasoactive intestinal peptide receptor 1 (VIPR1), mRNA
NM 030967	Homo sapiens keratin associated protein 1.1 (KRTAP1.1), mRNA
NM 015696	Homo sapiens weakly similar to glutathione peroxidase 2 (CL683), mRNA
	Homo sapiens Bardet-Biedl syndrome 2 (BBS2), mRNA
NM_031885	Homo sapiens keratin associated protein 1.3 (KRTAP1.3), mRNA
NM_030966	Homo sapiens nudix (nucleoside diphosphate linked moiety X)-type motif 6
NM_007083	
ND 4 012217	(NUDT6), mRNA
NM_013317	Homo sapiens lung type-I cell membrane-associated glycoprotein (T1A-2),
ND4 006474	transcript variant 1, mRNA
NM_006474	Homo sapiens lung type-I cell membrane-associated glycoprotein (T1A-2),
NR 6 00 6075	transcript variant 2, mRNA
NM_006275	Homo sapiens splicing factor, arginine/serine-rich 6 (SFRS6), mRNA
NM_016041	Homo sapiens CGI-101 protein (F-LAN-1), mRNA
NM_001954	Homo sapiens discoidin domain receptor family, member 1 (DDR1), transcript
	variant 2, mRNA
NM_013994	Homo sapiens discoidin domain receptor family, member 1 (DDR1), transcript
	variant 3, mRNA
NM_013993	Homo sapiens discoidin domain receptor family, member 1 (DDR1), transcript
	variant 1, mRNA
NM_022117	Homo sapiens cutaneous T-cell lymphoma-associated tumor antigen se20-4;
	differentially expressed nucleolar TGF-beta1 target protein (DENTT) (SE20-4),

	mRNA
NM_003048	Homo sapiens solute carrier family 9 (sodium/hydrogen exchanger), isoform 2 (SLC9A2), mRNA
NM 001971	Homo saniens elastase 1 nancreatic (ELA1), mRNA
NM 033412	Homo saniens hypothetical protein similar to CG/943 (MGC14836), mRNA
NM 033420	Homo saniens hypothetical protein MGC4022 (R32184_3), mRNA
NM 033408	Homo saniens hypothetical protein MBC3205 (MBC3205), mRNA
NM_014395	Homo sapiens dual adaptor of phosphotyrosine and 3-phosphoinositides
MM_014393	(DAPP1), mRNA
NM 003918	Homo saniens glycogenin 2 (GYG2), mRNA
NM 001502	Homo saniens glycoprotein 2 (zymogen granule memorane) (GP2), IIIXIA
NM 006362	Homo seniens nuclear RNA export factor 1 (NXF1), mRNA
NM 033155	Homo saniens nuclear RNA export factor 5 (NXF5), transcript variant 5, mkNA
NM 033154	Home seniers nuclear RNA export factor 5 (NXF5), transcript variant 4, HIKINA
NM 033153	Homo saniens nuclear RNA export factor 5 (NXF5), transcript variant 3, mRNA
NM 033152	Home capiens nuclear RNA export factor 5 (NXF5), transcript variant 2, mixiva
NM 032946	Homo sapiens nuclear RNA export factor 5 (NXF5), transcript variant 1, mRNA
NM 022052	Homo saniens nuclear RNA export factor 3 (NXF3), mRNA
NM 021808	Homo sapiens UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-
NWI_021808	acetylgalactosaminyltransferase 9 (GalNAc-19) (GALN19), mRNA
NM 017840	Homo sapiens mitochondrial ribosomal protein L16 (MRPL16), mRNA
NM 017417	Homo sapiens UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-
NWI_01/41/	acetylgalactosaminyltransferase 8 (GalNAc-T8) (GALNT8), mRNA
NM 004261	Homo saniens 15 kDa selenoprotein (SEP15), mRNA
NM 021998	Homo saniens zinc finger protein 6 (CMPX1) (ZNF6), mRNA
NM_004570	Homo sapiens phosphoinositide-3-kinase, class 2, gamma polypeptide
14141_004570	(DTV2C2C) mPNA
NM 002646	Homo sapiens phosphoinositide-3-kinase, class 2, beta polypeptide (PIK3C2B),
14141_002040	mPNA
NM 004598	Homo sapiens sparc/osteonectin, cwcv and kazal-like domains proteoglycan
14141_00 1550	(testionn) (SPOCK) mRNA
NM_033135	Homo sapiens spinal cord-derived growth factor-B (SCDGF-B), transcript
1111_000100	remient 2 mPNA
NM_025208	Homo sapiens spinal cord-derived growth factor-B (SCDGF-B), transcript
	variant 1 mRNA
NM 033346	Homo sapiens bone morphogenetic protein receptor, type II (serine/threonine
	lainage) (RMPR2) transcript variant 2, mRNA
NM 001204	Homo sapiens bone morphogenetic protein receptor, type II (serine/threonine
_	kinase) (BMPR2), transcript variant 1, mRNA
NM_003933	Homo saniens BAII-associated protein 3 (BAIAP3), mRNA
NM 005467	Homo sapiens N-acetylated alpha-linked acidic dipeptidase 2 (NAALAD2),
_	
NM_005944	Homo sapiens antigen identified by monoclonal antibody MRC OX-2 (MOX2),
1111_0000	DNIA
NM 002245	Homo sapiens potassium channel, subfamily K, member 1 (TWIK-1) (KCNK1),
	mRNA
NM 005247	Homo sapiens fibroblast growth factor 3 (murine mammary tumor virus
_	integration site (v-int-2) oncogene homolog) (FGF3), mRNA
NM 002006	Homo saniens fibroblast growth factor 2 (basic) (FGF2), mRNA
NM 000647	t i (a a .: \a
_	mDNA
NM 032047	Homo sapiens UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase

	5 (B3GNT5), mRNA
NIM 014256	Homo sapiens UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase
NM_014256	3 (B3GNT3), mRNA
NM 015904	Homo sapiens translation initiation factor IF2 (IF2), mRNA
NM_005326	* <del>                                     </del>
	Homo sapiens hydroxyacyl glutathione hydrolase (HAGH), mRNA
NM_013445	Homo sapiens glutamate decarboxylase 1 (brain, 67kD) (GAD1), transcript variant GAD25, mRNA
NM_033173	Homo sapiens UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide
	5 (B3GALT5), transcript variant 5, mRNA
NM 033172	Homo sapiens UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide
_	5 (B3GALT5), transcript variant 4, mRNA
NM_033171	Homo sapiens UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide
_	5 (B3GALT5), transcript variant 3, mRNA
NM_033170	Homo sapiens UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide
_	5 (B3GALT5), transcript variant 2, mRNA
NM_033169	Homo sapiens UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide
	3 (B3GALT3), transcript variant 4, mRNA
NM_033168	Homo sapiens UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide
	3 (B3GALT3), transcript variant 3, mRNA
NM_033167	Homo sapiens UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide
	3 (B3GALT3), transcript variant 2, mRNA
NM_003781	Homo sapiens UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide
	3 (B3GALT3), transcript variant 1, mRNA
NM_003782	Homo sapiens UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide
	4 (B3GALT4), mRNA
NM_003783	Homo sapiens UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide
	2 (B3GALT2), mRNA
NM_004631	Homo sapiens low density lipoprotein receptor-related protein 8, apolipoprotein
	e receptor (LRP8), transcript variant 1, mRNA
NM_033300	Homo sapiens low density lipoprotein receptor-related protein 8, apolipoprotein
	e receptor (LRP8), transcript variant 2, mRNA
NM_017522	Homo sapiens low density lipoprotein receptor-related protein 8, apolipoprotein
	e receptor (LRP8), transcript variant 3, mRNA
NM_033323	Homo sapiens sodium bicarbonate transporter 4 (NBC4), transcript variant b,
	mRNA
NM_033337	Homo sapiens caveolin 3 (CAV3), transcript variant 1, mRNA
NM_001234	Homo sapiens caveolin 3 (CAV3), transcript variant 2, mRNA
NM_001233	Homo sapiens caveolin 2 (CAV2), mRNA
NM_001753	Homo sapiens caveolin 1, caveolae protein, 22kD (CAV1), mRNA
NM_033291	Homo sapiens midline 1 (Opitz/BBB syndrome) (MID1), transcript variant 2,
	mRNA
NM_033290	Homo sapiens midline 1 (Opitz/BBB syndrome) (MID1), transcript variant 3,
	mRNA
NM_033274	Homo sapiens a disintegrin and metalloproteinase domain 19 (meltrin beta)
	(ADAM19), transcript variant 2, mRNA
NM_023038	Homo sapiens a disintegrin and metalloproteinase domain 19 (meltrin beta)
<del></del>	(ADAM19), transcript variant 1, mRNA
NM_033308	Homo sapiens ATP-binding cassette, sub-family A (ABC1), member 7
	(ABCA7), transcript variant 2, mRNA
NM_019112	Homo sapiens ATP-binding cassette, sub-family A (ABC1), member 7
37.6.000.000	(ABCA7), transcript variant 1, mRNA
NM_002609	Homo sapiens platelet-derived growth factor receptor, beta polypeptide

	CON CERTS PNA
	(PDGFRB), mRNA Homo sapiens platelet-derived growth factor receptor, alpha polypeptide
NM_006206	Homo sapiens platelet-derived growth factor receptor, diplica postpression
	(PDGFRA), mRNA  Homo sapiens platelet-derived growth factor beta polypeptide (simian sarcoma
NM_033016	viral (v-sis) oncogene homolog) (PDGFB), transcript variant 2, mRNA
	Homo sapiens adrenergic, alpha-1D-, receptor (ADRA1D), mRNA
NM_000678	Homo sapiens adrenergic, alpha-1B-, receptor (ADRA1B), mRNA  Homo sapiens adrenergic, alpha-1B-, receptor (ADRA1B), mRNA
NM_000679	Homo sapiens promyelocytic leukemia (PML), transcript variant 6, mRNA
NM_002675	Homo sapiens promyelocytic leukemia (PML), transcript variant 11, mRNA  Homo sapiens promyelocytic leukemia (PML), transcript variant 11, mRNA
NM_033250	Homo sapiens promyelocytic leukemia (PML), transcript variant 10, mRNA
NM_033249	Homo sapiens promyelocytic leukemia (PML), transcript variant 8, mRNA  Homo sapiens promyelocytic leukemia (PML), transcript variant 8, mRNA
NM_033247	Homo sapiens promyelocytic leukemia (PML), transcript variant 7, mRNA
NM_033246	Homo sapiens promyelocytic leukemia (PML), transcript variant 1, mRNA  Homo sapiens promyelocytic leukemia (PML), transcript variant 12, mRNA
NM_033245	Homo sapiens promyelocytic leukeinia (FWL), transcript variant 5 mRNA
NM_033244	Homo sapiens promyelocytic leukemia (PML), transcript variant 5, mRNA
NM_033242	Homo sapiens promyelocytic leukemia (PML), transcript variant 3, mRNA
NM_033240	Homo sapiens promyelocytic leukemia (PML), transcript variant 2, mRNA
NM_033239	Homo sapiens promyelocytic leukemia (PML), transcript variant 9, mRNA
NM_033238	Homo sapiens promyelocytic leukemia (PML), transcript variant 1, mRNA
NM_033304	Homo sapiens adrenergic, alpha-1A-, receptor (ADRA1A), transcript variant 4, mRNA
NM_033303	Homo sapiens adrenergic, alpha-1A-, receptor (ADRA1A), transcript variant 2, mRNA
NM_033302	Homo sapiens adrenergic, alpha-1A-, receptor (ADRA1A), transcript variant 3,
NM 033279	Homo saniens ring finger protein 22 (RNF22), transcript variant gamma, mRNA
NM 033278	Homo saniens ring finger protein 22 (RNF22), transcript variant beta, mkNA
NM 000737	Homo saniens charionic gonadotropin, beta polypeptide (CGB), mRNA
NM 033295	Homo sapiens caspase 1, apoptosis-related cysteine protease (interleukin 1, beta,
14141_033293	convertees) (CASP1) transcript variant ensilon, mRNA,
NM_033294	Homo sapiens caspase 1, apoptosis-related cysteine protease (interleukin 1, beta,
14141_055254	convertage) (CASP1) transcript variant delta, mKNA
NM_033293	Homo sapiens caspase 1, apoptosis-related cysteine protease (interleukin 1, beta,
14141_053253	convertase) (CASP1) transcript variant gamma, mRNA
NM_033292	Homo sapiens caspase 1, apoptosis-related cysteine protease (interleukin 1, beta, convertase) (CASPI) transcript variant alpha, mRNA
ND4 001222	Homo sapiens caspase 1, apoptosis-related cysteine protease (interleukin 1, beta,
NM_001223	convertase) (CASP1), transcript variant beta, mRNA
ND ( 006771	Homo sapiens keratin, hair, acidic, 8 (KRTHA8), mRNA
NM_006771	Homo sapiens keratin, hair, acidic, 5 (KRTHA5), mRNA
NM_002280	Homo sapiens keratin 14 (epidermolysis bullosa simplex, Dowling-Meara,
NM_000526	Koebner) (KRT14), mRNA
ND 6 022201	Homo sapiens ribosomal protein L8 (RPL8), transcript variant 2, mRNA
NM_033301	Homo sapiens ribosomal protein L8 (RPL8), transcript variant 1, mRNA
NM_000973	Homo sapiens ribosomal protein L9 (RPL9), mRNA
NM_000661	Homo saniens ribosomal protein L10a (RPL10A), mRNA
NM_007104	Homo saniens enithelial stromal interaction I (breast) (EPSIII), mRNA
NM 033255	Homo sapiens sodium bicarbonate transporter 4 (NBC4), transcript variant a,
NM_021196	mRNA
NM_032241	Homo sapiens ribosomal protein L10 (RPL10), mRNA
NM_030955	Homo sapiens a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 12 (ADAMTS12), mRNA
NM_030765	Homo sapiens UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase

	4 (D2CNT4)DNA
27.6.01.4670	4 (B3GNT4), mRNA
NM_014670	Homo sapiens basic leucine-zipper protein BZAP45 (BZAP45), mRNA
NM_013379	Homo sapiens dipeptidylpeptidase 7 (DPP7), mRNA
NM_006458	Homo sapiens ring finger protein 22 (RNF22), transcript variant alpha, mRNA
NM_006057	Homo sapiens UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide
37.6.000640	5 (B3GALT5), transcript variant 1, mRNA
NM_000648	Homo sapiens chemokine (C-C motif) receptor 2 (CCR2), transcript variant B, mRNA
NM_000381	Homo sapiens midline 1 (Opitz/BBB syndrome) (MID1), transcript variant 1, mRNA
NM_002645	Homo sapiens phosphoinositide-3-kinase, class 2, alpha polypeptide (PIK3C2A), mRNA
NM_002608	Homo sapiens platelet-derived growth factor beta polypeptide (simian sarcoma
11	viral (v-sis) oncogene homolog) (PDGFB), transcript variant 1, mRNA
NM 001134	Homo sapiens alpha-fetoprotein (AFP), mRNA
NM_000680	Homo sapiens adrenergic, alpha-1A-, receptor (ADRA1A), transcript variant 1, mRNA
NM 023929	Homo sapiens zinc finger protein RINZF (RINZF), mRNA
NM 020353	Homo sapiens phospholipid scramblase 4 (PLSCR4), mRNA
NM 020359	Homo sapiens phospholipid scramblase 2 (PLSCR2), mRNA
NM 018494	Homo sapiens leucine-rich and death domain containing (LRDD), mRNA
NM 004998	Homo sapiens myosin IE (MYO1E), mRNA
NM 033226	Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 12
	(ABCC12), mRNA
NM_032105	Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 12B (PPP1R12B), transcript variant 2, mRNA
NM_032104	Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 12B (PPP1R12B), transcript variant 4, mRNA
NM_032103	Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 12B (PPP1R12B), transcript variant 3, mRNA
NM_002481	Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 12B (PPP1R12B), transcript variant 1, mRNA
NM 004689	Homo sapiens metastasis associated 1 (MTA1), mRNA
NM 006005	Homo sapiens Wolfram syndrome 1 (wolframin) (WFS1), mRNA
NM_015722	Homo sapiens calcyon; D1 dopamine receptor-interacting protein (CALCYON), mRNA
NM 004184	Homo sapiens tryptophanyl-tRNA synthetase (WARS), mRNA
NM_014228	Homo sapiens solute carrier family 6 (neurotransmitter transporter, L-proline), member 7 (SLC6A7), mRNA
NM 005823	Homo sapiens mesothelin (MSLN), transcript variant 1, mRNA
NM 013404	Homo sapiens mesothelin (MSLN), transcript variant 1, mRNA
NM 012341	Homo sapiens G protein-binding protein CRFG (CRFG), mRNA
NM 002480	Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 12A
	(PPP1R12A), mRNA
NM_003868	Homo sapiens fibroblast growth factor 16 (FGF16), mRNA
NM_018979	Homo sapiens protein kinase, lysine deficient 1 (PRKWNK1), mRNA
NM_022127	Homo sapiens solute carrier family 28 (sodium-coupled nucleoside transporter), member 3 (SLC28A3), mRNA
NM_005517	Homo sapiens high-mobility group (nonhistone chromosomal) protein 17 (HMG17), mRNA
NM 022465	Homo sapiens zinc finger protein, subfamily 1A, 4 (Eos) (ZNFN1A4), mRNA
NM 005768	Homo sapiens putative protein similar to nessy (Drosophila) (C3F), mRNA
2111 003700	1 Home supno putting butter to heady (151000pmin) (Cot ), Illicant

Homo sapiens stresscopin-related peptide (SRP), mRNA
Homo sapiens thioredoxin domain-containing 2 (spermatozoa) (TXNDC2),
mRNA (ACED D) DNA
Homo sapiens membrane-type frizzled-related protein (MFRP), mRNA
Homo sapiens zinc finger protein, subfamily 1A, 5 (Pegasus) (PEGASUS),
mRNA
Homo sapiens ATPase, Ca++ transporting, cardiac muscle, fast twitch 1
(ATP2A1), mRNA
Homo sapiens zinc finger protein 253 (ZNF253), mRNA
Homo sapiens chromosome 21 open reading frame 7 (C21orf7), mRNA
Homo sapiens chromosome 21 open reading frame 91 (C21orf91), mRNA  Homo sapiens chromosome 21 open reading frame 91 (C21orf91), mRNA
Homo sapiens RAB4B, member RAS oncogene family (RAB4B), mRNA
Homo sapiens UMP-CMP kinase (UMP-CMPK), mRNA
Homo sapiens glutaredoxin 2 (GLRX2), mRNA
Homo sapiens family with sequence similarity 8, member A1 (FAM8A1),
mRNA
Homo sapiens likely ortholog of mouse coiled coil forming protein 1
(KIAA0203), mRNA
Homo sapiens VENT-like homeobox 2 (VENTX2), mRNA
Homo sapiens transcription factor-like 4 (TCFL4), mRNA
Homo sapiens zinc finger protein, subfamily 1A, 3 (Aiolos) (ZNFN1A3), mRNA  Homo sapiens zinc finger protein, subfamily 1A, 3 (Aiolos) (ZNFN1A3), mRNA
Homo sapiens POM (POM121 rat homolog) and ZP3 fusion (POMZP3), mRNA
Homo sapiens eukaryotic translation initiation factor 2C, 1 (EIF2C1), mRNA
Homo sapiens immunoglobulin superfamily, member 6 (IGSF6), mRNA
Homo sapiens SKI-like (SKIL), mRNA
Homo sapiens transglutaminase 5 (TGM5), mRNA
Homo sapiens megakaryoblastic leukemia (translocation) 1 (MKL1), mRNA
Homo sapiens endogenous retrovirus H D1 leader region/integrase-derived
ORF1, ORF2, and putative envelope protein (HSU88895), mRNA
Homo sapiens scavenger receptor cysteine-rich type 1 protein M160 precursor
(M160), mRNA
Homo sapiens Sox-6 (HSSOX6), mRNA Homo sapiens cat eye syndrome chromosome region, candidate 5 (CECR5),
Homo sapiens cat eye syndrome chromosome region, candidate s
mRNA Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 14A
Homo sapiens protein phosphatase 1, regulatory (minorot) see that a
(PPP1R14A), mRNA  Homo sapiens hypothetical protein MGC12466 (MGC12466), mRNA
Homo sapiens nypothetical protein MGC12400 (MGC12400),  Homo sapiens cat eye syndrome chromosome region, candidate 5 (CECR5),
mRNA Homo sapiens hypothetical protein MGC15548 (MGC15548), mRNA
Homo sapiens hypothetical protein MGC13008 (MGC13008), mRNA
Homo sapiens hypothetical protein MGC15416 (MGC15416), mRNA
Homo sapiens hypothetical protein MGC13114 (MGC13114), mRNA  Homo sapiens hypothetical protein MGC13114 (MGC13114), mRNA
Homo sapiens hypothetical protein MGC10540 (MGC10540), mRNA
Homo sapiens hypothetical protein MGC2605 (MGC2605), mRNA
Homo sapiens hypothetical protein MGC2003 (MGC2003), max 12 Homo sapiens hypothetical protein DKFZp434F054 (DKFZp434F054), mRNA
Homo sapiens hypothetical protein FLJ23519 (FLJ23519), mRNA
Homo sapiens hypothetical protein FL323319 (FL323319), inicity  Homo sapiens zinc family member 4 protein HZIC4 (ZIC4), mRNA
Homo sapiens zinc family incliner 4 protein fizzio (2104), inclui
Homo sapiens ELKS protein (ELKS), mRNA Homo sapiens hypothetical protein DKFZp586M1120 (DKFZP586M1120),
mRNA

NM_025267	Homo sapiens hypothetical protein MGC2744 (MGC2744), mRNA
NM_025051	Homo sapiens hypothetical protein FLJ23022 (FLJ23022), mRNA
NM_024974	Homo sapiens hypothetical protein FLJ11800 (FLJ11800), mRNA
NM_024934	Homo sapiens hypothetical protein FLJ22659 (FLJ22659), mRNA
NM 024805	Homo sapiens hypothetical protein FLJ21172 (FLJ21172), mRNA
NM 024804	Homo sapiens hypothetical protein FLJ12606 (FLJ12606), mRNA
NM 024052	Homo sapiens hypothetical protein MGC3048 (MGC3048), mRNA
NM 024042	Homo sapiens hypothetical protein MGC2601 (MGC2601), mRNA
NM_020535	Homo sapiens killer cell immunoglobulin-like receptor, two domains, long
_	cytoplasmic tail, 5 (KIR2DL5), mRNA
NM_021939	Homo sapiens hypothetical protein FLJ22041 similar to FK506 binding proteins
	(FLJ22041), mRNA
NM_020664	Homo sapiens 2,4-dienoyl CoA reductase 2, peroxisomal (DECR2), mRNA
NM_018722	Homo sapiens BWRT protein (HSA404617), mRNA
NM_020394	Homo sapiens zinc finger protein SBZF3 (LOC57116), mRNA
NM_019013	Homo sapiens hypothetical protein (FLJ10156), mRNA
NM 018629	Homo sapiens hypothetical protein PRO2533 (PRO2533), mRNA
NM 018568	Homo sapiens hypothetical protein PRO0943 (PRO0943), mRNA
NM 018050	Homo sapiens hypothetical protein FLJ10298 (FLJ10298), mRNA
NM 018019	Homo sapiens hypothetical protein FLJ10193 (FLJ10193), mRNA
NM 017609	Homo sapiens hypothetical protein DKFZp434A1721 (DKFZp434A1721),
_	mRNA
NM_016332	Homo sapiens selenoprotein X, 1 (SEPX1), mRNA
NM 016360	Homo sapiens clone HQ0477 PRO0477p (LOC51204), mRNA
NM 016002	Homo sapiens CGI-49 protein (LOC51097), mRNA
NM 014913	Homo sapiens KIAA0863 protein (KIAA0863), mRNA
NM 014700	Homo sapiens KIAA0665 gene product (KIAA0665), mRNA
NM 014680	Homo sapiens KIAA0100 gene product (KIAA0100), mRNA
NM 012248	Homo sapiens selenophosphate synthetase 2 (SPS2), mRNA
NM 007222	Homo sapiens zinc-fingers and homeoboxes 1 (ZHX1), mRNA
NM 006555	Homo sapiens SNARE protein (YKT6), mRNA
NM_006623	Homo sapiens phosphoglycerate dehydrogenase (PHGDH), mRNA
NM 006613	Homo sapiens GRB2-related adaptor protein (GRAP), mRNA
NM 006659	Homo sapiens gamma-tubulin complex protein 2 (GCP2), mRNA
NM 016441	Homo sapiens cysteine-rich motor neuron 1 (CRIM1), mRNA
NM_014787	Homo sapiens DnaJ (Hsp40) homolog, subfamily C, member 6 (DNAJC6), mRNA
NM_004213	Homo sapiens solute carrier family 28 (sodium-coupled nucleoside transporter), member 1 (SLC28A1), mRNA
NM_003141	Homo sapiens Sjogren syndrome antigen A1 (52kD, ribonucleoprotein autoantigen SS-A/Ro) (SSA1), mRNA
NM_002607	Homo sapiens platelet-derived growth factor alpha polypeptide (PDGFA), transcript variant 1, mRNA
NM_033023	Homo sapiens platelet-derived growth factor alpha polypeptide (PDGFA), transcript variant 2, mRNA
NM 005675	Homo sapiens DiGeorge syndrome critical region gene 6 (DGCR6), mRNA
NM_016083	Homo sapiens cannabinoid receptor 1 (brain) (CNR1), transcript variant 2, mRNA
NM 004053	Homo sapiens bystin-like (BYSL), mRNA
NG_000016	Homo sapiens genomic protocadherin alpha cluster (PCDHA@) on chromosome 5
NM 032935	Homo sapiens metallothionein IV (MTIV), mRNA

	D (D40)DNA
NM_003695	Homo sapiens lymphocyte antigen 6 complex, locus D (E48), mRNA
NM_006787	Homo sapiens melanoma antigen, family D, 2 (MAGED2), mRNA
NM_016205	Homo sapiens platelet derived growth factor C (PDGFC), mRNA
NM_017913	Homo sapiens Hsp90-associating relative of Cdc37 (HARC), mRNA
VM_017701	Homo sapiens Rho GTPase activating protein 8 (ARHGAP8), mRNA
VM_015366	Homo sapiens Rho GTPase activating protein 8 (ARHGAP8), mRNA
NM_012269	Homo sapiens hyaluronoglucosaminidase 4 (HYAL4), mRNA
NM_006207	Homo sapiens platelet-derived growth factor receptor-like (PDGFRL), mRNA
NM_004986	Homo sapiens kinectin 1 (kinesin receptor) (KTN1), mRNA
NM_001840	Homo sapiens cannabinoid receptor 1 (brain) (CNR1), transcript variant 1, mRNA
NM_014417	Homo saniens Bcl-2 binding component 3 (BBC3), mRNA
NM_033223	Homo sapiens gamma-aminobutyric acid (GABA) A receptor, gamma 3 (GABRG3), mRNA
NM_005762	Homo saniens tripartite motif-containing 28 (TRIM28), mRNA
NM_015906	Homo sapiens tripartite motif-containing 33 (TRIM33), transcript variant aipna,
NM_033020	Homo sapiens tripartite motif-containing 33 (TRIM33), transcript variant beta,
NM 032421	Homo saniens cytoplasmic linker 2 (CYLN2), transcript variant 2, mRNA
NM 031416	Homo sapiens chromosome 18 open reading frame 2 (C18orf2), mRNA
NM 014412	Homo sapiens Siah-interacting protein (SIP), mRNA
NM 016212	Homo sapiens TP53TG3 protein (TP53TG3), mRNA
NM 016552	Homo sapiens testis specific ankyrin-like protein 1 (LOC51281), mRNA
NM 015369	Homo sapiens TP53TG3 protein (TP53TG3), mRNA
NM 033284	Homo sapiens transducin beta-like 1 protein (TBL1Y), mRNA
NM 031951	Homo sapiens NYD-SP11 protein (NYD-SP11), mRNA
NM_020414	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 24 (DDX24), mRNA
NM 007268	Homo sapiens Ig superfamily protein (Z39IG), mRNA
NM 006707	Homo saniens butyrophilin-like 3 (BTNL3), mRNA
NM_002491	Homo sapiens NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 3 (12kD, B12) (NDUFB3), mRNA
NM_001386	Homo sapiens dihydropyrimidinase-like 2 (DPYSL2), mRNA
NM_000090	Homo sapiens collagen, type III, alpha 1 (Ehlers-Danlos syndrome type IV, autosomal dominant) (COL3A1), mRNA
NM_033150	Homo sapiens collagen, type II, alpha 1 (primary osteoarthritis, spondyloepiphyseal dysplasia, congenital) (COL2A1), transcript variant 2, mRNA
NM_001844	Homo sapiens collagen, type II, alpha 1 (primary osteoarthritis, spondyloepiphyseal dysplasia, congenital) (COL2A1), transcript variant 1, mRNA
NM_025245	Homo sapiens pre-B-cell leukemia transcription factor 4 (PBX4), mRNA
NM_004342	Homo sapiens caldesmon 1 (CALD1), transcript variant 3, mRNA
NM_033157	Homo sapiens caldesmon 1 (CALD1), transcript variant 2, mRNA
NM_033140	Homo sapiens caldesmon 1 (CALD1), transcript variant 5, mRNA
NM_033139	Homo sapiens caldesmon 1 (CALD1), transcript variant 4, mRNA
NM 033138	Homo saniens caldesmon 1 (CALD1), transcript variant 1, mRNA
NM 032635	Homo sapiens seven transmembrane domain protein (NIFIE14), mRNA
NM 030912	Homo sapiens ring finger protein 27 (RNF27), mRNA
	Homo sapiens solute carrier family 7, (cationic amino acid transporter, y+

NM 017844	Homo sapiens testis specific ankyrin-like protein 1 (LOC51281), mRNA
NM 014242	Homo sapiens zinc finger protein 237 (ZNF237), mRNA
NM 001715	Homo sapiens B lymphoid tyrosine kinase (BLK), mRNA
NM 033158	Homo sapiens hyaluronoglucosaminidase 2 (HYAL2), transcript variant 2,
-	mRNA
NM 033159	Homo sapiens hyaluronoglucosaminidase 1 (HYAL1), transcript variant 2,
_	mRNA
NM_007312	Homo sapiens hyaluronoglucosaminidase 1 (HYAL1), transcript variant 1,
	mRNA
NM_006119	Homo sapiens fibroblast growth factor 8 (androgen-induced) (FGF8), transcript
	variant B, mRNA
NM_033165	Homo sapiens fibroblast growth factor 8 (androgen-induced) (FGF8), transcript
	variant A, mRNA
NM_033164	Homo sapiens fibroblast growth factor 8 (androgen-induced) (FGF8), transcript
	variant E, mRNA
NM_033163	Homo sapiens fibroblast growth factor 8 (androgen-induced) (FGF8), transcript
	variant F, mRNA
NM_002009	Homo sapiens fibroblast growth factor 7 (keratinocyte growth factor) (FGF7),
27 6 001005	mRNA
NM_021907	Homo sapiens dystrobrevin, beta (DTNB), transcript variant 1, mRNA
NM_033148	Homo sapiens dystrobrevin, beta (DTNB), transcript variant 3, mRNA
NM_033147	Homo sapiens dystrobrevin, beta (DTNB), transcript variant 2, mRNA
NM_015902	Homo sapiens progestin induced protein (DD5), mRNA
NM_000777	Homo sapiens cytochrome P450, subfamily IIIA (niphedipine oxidase),
ND 6 000764	polypeptide 5 (CYP3A5), mRNA  Homo sapiens cytochrome P450, subfamily IIA (phenobarbital-inducible),
NM_000764	polypeptide 7 (CYP2A7), transcript variant 1, mRNA
NM 030589	Homo sapiens cytochrome P450, subfamily IIA (phenobarbital-inducible),
MM_030389	polypeptide 7 (CYP2A7), transcript variant 2, mRNA
NM 000762	Homo sapiens cytochrome P450, subfamily IIA (phenobarbital-inducible),
14141_000702	polypeptide 6 (CYP2A6), mRNA
NM 018957	Homo sapiens SH3-domain binding protein 1 (SH3BP1), mRNA
NM 033258	Homo sapiens G-protein gamma 8 subunit (GNG8), mRNA
NM 033260	Homo sapiens winged helix/forkhead transcription factor (HFH1), mRNA
NM 018476	Homo sapiens brain expressed, X-linked 1 (BEX1), mRNA
NM 022154	Homo sapiens up-regulated by BCG-CWS (LOC64116), mRNA
NM 003773	Homo sapiens hyaluronoglucosaminidase 2 (HYAL2), transcript variant 1,
	mRNA
NM 032794	Homo sapiens NG22 protein (NG22), mRNA
NM_030768	Homo sapiens integrin-linked kinase-associated serine/threonine phosphatase 2C
	(ILKAP), mRNA
NM_025257	Homo sapiens NG22 protein (NG22), mRNA
NM_020996	Homo sapiens fibroblast growth factor 6 (FGF6), mRNA
NM_016543	Homo sapiens sialic acid binding Ig-like lectin 7 (SIGLEC7), mRNA
NM_016134	Homo sapiens plasma glutamate carboxypeptidase (PGCP), mRNA
NM 014385	Homo sapiens sialic acid binding Ig-like lectin 7 (SIGLEC7), mRNA
NM_013287	Homo sapiens phosphoprotein enriched in astrocytes 15 (PEA15), mRNA
NM_006102	Homo sapiens plasma glutamate carboxypeptidase (PGCP), mRNA
NM_004112	Homo sapiens fibroblast growth factor 11 (FGF11), mRNA
NM_004465	Homo sapiens fibroblast growth factor 10 (FGF10), mRNA
NM_003811	Homo sapiens tumor necrosis factor (ligand) superfamily, member 9 (TNFSF9),
	mRNA

T	VY CONTRACTOR (CLND mPNA
NM_003063	Homo sapiens sarcolipin (SLN), mRNA
NM_003768	Homo sapiens phosphoprotein enriched in astrocytes 15 (PEA15), mRNA
NM_002010	Homo sapiens fibroblast growth factor 9 (glia-activating factor) (FGF9), mRNA
NM_033215	Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 3F
	(PPP1R3F), mRNA
NM_032741	Homo sapiens 1-acylglycerol-3-phosphate O-acyltransferase 1 (lysophosphatidic
_	acid acyltransferase, alpha) (AGPAT1), mRNA
NM 022152	Homo sapiens PP1201 protein (PP1201), mRNA
NM 033225	Homo sapiens CUB and Sushi multiple domains 1 (CSMD1), mRNA
NM 014505	Homo sapiens potassium large conductance calcium-activated channel,
	subfamily M, beta member 4 (KCNMB4), mRNA
NM 032559	Homo sapiens kinesin protein (LOC84643), mRNA
NM 015394	Homo sapiens zinc finger protein 10 (KOX 1) (ZNF10), mRNA
NM 003388	Homo saniens cytoplasmic linker 2 (CYLN2), transcript variant 1, mRNA
NM 032736	Homo sapiens torsin family 1, member B (torsin B) (TOR1B), mRNA
NM 032689	Homo sapiens hypothetical protein MGC13071 (MGC13071), mRNA
NM 032227	Homo saniens hypothetical protein FLJ22679 (FLJ22679), mRNA
NM 014506	Homo sapiens torsin family 1, member B (torsin B) (TOR1B), mRNA
NM 030900	Homo sapiens cell cycle progression 2 protein (CPR2), mRNA
	Homo sapiens oxysterol binding protein 2 (OSBP2), mRNA
NM_030758	Homo sapiens hypothetical protein FLJ22679 (FLJ22679), mRNA
NM_017698	Homo sapiens homolog of C. elegans smu-1 (SMU-1), mRNA
NM_018225	Homo sapiens Kruppel-like factor 12 (KLF12), mRNA
NM_016285	Homo sapiens Kruppel-like factor 12 (KLF12), mRNA
NM_007249	Homo sapiens Kruppel-like factor 12 (KLF12), mRNA
NM_006464	Homo sapiens trans-golgi network protein 2 (TGOLN2), mRNA
NM_006411	Homo sapiens 1-acylglycerol-3-phosphate O-acyltransferase 1 (lysophosphatidic
	acid acyltransferase, alpha) (AGPAT1), mRNA
NM_004749	Homo sapiens cell cycle progression 2 protein (CPR2), mRNA
NM_000285	Homo sapiens peptidase D (PEPD), mRNA
NM_001467	Homo sapiens glucose-6-phosphatase, transport (glucose-6-phosphate) protein 1
37 f 022108	(G6PT1), mRNA  Homo sapiens phosphatidylinositol glycan, class S (PIGS), mRNA
NM_033198	Homo sapiens phosphatidylmoshor gyean, class 8 (1905), me in Homo sapiens regulatory factor X, 4 (influences HLA class II expression)
NM_002920	
	(RFX4), mRNA
NM_018944	Homo sapiens chromosome 21 open reading frame 45 (C21orf45), mRNA
NM_033214	Homo sapiens glycerol kinase pseudogene 2 (GKP2), mRNA
NM_033089	Homo sapiens hypothetical protein FLJ22115 (FLJ22115), mRNA
NM_016015	Homo sapiens leucine carboxyl methyltransferase (LCMT), mRNA
NM_033209	Homo sapiens Thy-1 co-transcribed (LOC94105), mRNA
NM_033093	Homo sapiens tripartite motif-containing 5 (TRIM5), transcript variant delta,
	mRNA
NM_033092	Homo sapiens tripartite motif-containing 5 (TRIM5), transcript variant gamma,
	mRNA
NM_033091	Homo sapiens tripartite motif-containing 4 (TRIM4), transcript variant beta,
	mRNA (TDD 14)
NM_033017	Homo sapiens tripartite motif-containing 4 (TRIM4), transcript variant alpha,
	mRNA (TD P (S)
NM_033034	Homo sapiens tripartite motif-containing 5 (TRIM5), transcript variant alpha,
_	mRNA
NM_015318	Homo sapiens Rho-specific guanine nucleotide exchange factor p114 (P114-
	RHO-GEF), mRNA
NM 007204	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 20, 103kD

	(DDV20) DNA
ND4 022864	(DDX20), mRNA  Homo sapiens hypothetical protein FLJ14936 (FLJ14936), mRNA
NM_032864 NM_032639	Homo sapiens phosphoinositol 4-phosphate adaptor protein-2 (FAPP2), mRNA
	Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 11
NM_032583	(ABCC11), mRNA
NIM 022284	Homo sapiens hypothetical protein FLJ14936 (FLJ14936), mRNA
NM_032284	Homo sapiens hypothetical protein FLJ13614 (FLJ13614), mRNA
NM_032182	Homo sapiens fatty acid desaturase 3 (FADS3), mRNA
NM_021727 NM_022726	Homo sapiens elongation of very long chain fatty acids (FEN1/Elo2, SUR4/Elo3,
NNI_022720	yeast)-like 4 (ELOVL4), mRNA
NM_015162	Homo sapiens lipidosin (BG1), mRNA
NM_021176	Homo sapiens islet-specific glucose-6-phosphatase catalytic subunit-related protein (IGRP), mRNA
NM_019094	Homo sapiens nudix (nucleoside diphosphate linked moiety X)-type motif 4 (NUDT4), mRNA
NM_019091	Homo sapiens pleckstrin homology domain-containing, family A
1111_017071	(phosphoinositide binding specific) member 3 (PLEKHA3), mRNA
NM 018293	Homo sapiens hypothetical protein FLJ10997 (FLJ10997), mRNA
NM 015994	Homo sapiens ATPase, H+ transporting lysosomal (vacuolar proton pump),
	member M (ATP6M), mRNA
NM 015952	Homo sapiens PTD013 protein (PTD013), mRNA
NM 015899	Homo sapiens putative glycolipid transfer protein (LOC51054), mRNA
NM 016309	Homo sapiens leucine carboxyl methyltransferase (LCMT), mRNA
NM 013345	Homo sapiens G protein-coupled receptor (G2A), mRNA
NM 012228	Homo sapiens pilin-like transcription factor (PILB), mRNA
NM_006886	Homo sapiens ATP synthase, H+ transporting, mitochondrial F1 complex,
	epsilon subunit (ATP5E), mRNA
NM_002200	Homo sapiens interferon regulatory factor 5 (IRF5), transcript variant 1, mRNA
NM_032643	Homo sapiens interferon regulatory factor 5 (IRF5), transcript variant 2, mRNA
NM_004464	Homo sapiens fibroblast growth factor 5 (FGF5), transcript variant 1, mRNA
NM_033143	Homo sapiens fibroblast growth factor 5 (FGF5), transcript variant 2, mRNA
NM_020638	Homo sapiens fibroblast growth factor 23 (FGF23), mRNA
NM_000800	Homo sapiens fibroblast growth factor 1 (acidic) (FGF1), transcript variant 1, mRNA
NM_033137	Homo sapiens fibroblast growth factor 1 (acidic) (FGF1), transcript variant 3, mRNA
NM 032102	Homo sapiens Splicing factor, arginine/serine-rich, 46kD (SRP46), mRNA
NM_033136	Homo sapiens fibroblast growth factor 1 (acidic) (FGF1), transcript variant 2, mRNA
NM 002952	Homo sapiens ribosomal protein S2 (RPS2), mRNA
NM 033130	Homo sapiens sialic acid binding Ig-like lectin 10 (SIGLEC10), mRNA
NM 020665	Homo sapiens kidney-specific membrane protein (NX-17), mRNA
NM_033180	Homo sapiens olfactory receptor, family 51, subfamily B, member 2 (OR51B2), mRNA
NM_033179	Homo sapiens olfactory receptor, family 51, subfamily B, member 4 (OR51B4),
NIM 022179	mRNA Home services double homeobox 4 (DUV4) mRNA
NM_033178	Homo sapiens double homeobox, 4 (DUX4), mRNA
NM 033049	Homo sapiens mucin 13, epithelial transmembrane (MUC13), mRNA
NM_021619	Homo sapiens PR domain containing 12 (PRDM12), mRNA
NM_020382	Homo sapiens PR/SET domain containing protein 07 (SET07), mRNA
NM_007365	Homo sapiens peptidyl arginine deiminase, type II (PDI2), mRNA
NM_015894	Homo sapiens stathmin-like 3 (STMN3), mRNA

NIM 022401	Homo sapiens regulatory factor X, 4 (influences HLA class II expression)
NM_032491	(RFX4), mRNA
NM 024551	Homo sapiens hypothetical protein FLJ21432 (FLJ21432), mRNA
NM 021830	Homo sapiens chromosome 10 open reading frame 2 (C10orf2), mRNA
NM 017972	Homo sapiens hypothetical protein FLJ20689 (FLJ20689), mRNA
NM 020398	Homo sapiens serine protease inhibitor-like, with Kunitz and WAP domains 1
1411_020370	(eppin) (SPINLW1), mRNA
NM 020637	Homo sapiens fibroblast growth factor 22 (FGF22), mRNA
NM_019113	Homo sapiens fibroblast growth factor 21 (FGF21), mRNA
NM 017926	Homo sapiens hypothetical protein FLJ20689 (FLJ20689), mRNA
	Homo sapiens zinc finger protein 226 (ZNF226), mRNA
NM_016444	Homo sapiens serologically defined breast cancer antigen 84 (SDBCAG84),
NM_015966	mRNA
NM_015919	Homo sapiens zinc finger protein 226 (ZNF226), mRNA
NM 015474	Homo sapiens SAM domain and HD domain, 1 (SAMHD1), mRNA
NM_007096	Homo sapiens clathrin, light polypeptide (Lca) (CLTA), transcript variant brain- specific, mRNA
NM_002007	Homo sapiens fibroblast growth factor 4 (heparin secretory transforming protein
NM_002007	1, Kaposi sarcoma oncogene) (FGF4), mRNA
ND 6 001922	Homo sapiens clathrin, light polypeptide (Lca) (CLTA), transcript variant
NM_001833	
NR 6 000140	nonbrain, mRNA
NM_022143	Homo sapiens NAG14 protein (NAG14), mRNA
NM_005292	Homo sapiens G protein-coupled receptor 18 (GPR18), mRNA
NM_001371	Homo sapiens dynein, axonemal, heavy polypeptide 8 (DNAH8), mRNA
NM_012276	Homo sapiens leukocyte immunoglobulin-like receptor, subfamily A (without
	TM domain), member 4 (ILT7), mRNA
NM_012092	Homo sapiens inducible T-cell co-stimulator (ICOS), mRNA
NM_032447	Homo sapiens fibrillin3 (KIAA1776), mRNA
NM_024017	Homo sapiens homeo box B9 (HOXB9), mRNA
NM_019558	Homo sapiens homeo box D8 (HOXD8), mRNA
NM_032379	Homo sapiens synaptotagmin-like 2 (SYTL2), transcript variant b, mRNA
NM_024690	Homo sapiens mucin 16 (MUC16), mRNA
NM_018558	Homo sapiens gamma-aminobutyric acid (GABA) receptor, theta (GABRQ), mRNA
ND 6 014452	Homo sapiens tumor necrosis factor receptor superfamily, member 21
NM_014452	
ND 6 00 (042	(TNFRSF21), mRNA  Homo sapiens protein phosphatase 1, regulatory subunit 3D (PPP1R3D), mRNA
NM_006242	Homo sapiens homologous to yeast nitrogen permease (candidate tumor
NM_006545	
>7 6 005200	suppressor) (NPR2L), mRNA  Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 3C
NM_005398	Homo sapiens protein phosphatase 1, regulatory (minotor) subunit 30
	(PPP1R3C), mRNA  Homo sapiens serologically defined colon cancer antigen 28 (SDCCAG28),
NM_006645	Homo sapiens serologically defined colon cancer antigen 28 (SDCCAC28),
	mRNA
NM_032800	Homo sapiens hypothetical protein FLJ14525 (FLJ14525), mRNA
NM_004265	Homo sapiens fatty acid desaturase 2 (FADS2), mRNA
NM_013402	Homo sapiens fatty acid desaturase 1 (FADS1), mRNA
NM_031428	Homo sapiens hypothetical protein FLJ14525 (FLJ14525), mRNA
NM_025243	Homo sapiens solute carrier family 19, member 3 (SLC19A3), mRNA
NM 024411	Homo sapiens prodynorphin (PDYN), mRNA
NM 007368	Homo sapiens RAS p21 protein activator (GTPase activating protein) 3
_	(Ins(1,3,4,5)P4-binding protein) (GAP1IP4BP), mRNA
NM 003912	Homo sapiens myotubularin related protein 2 (MTMR2), mRNA

Homo sapiens ubiquitin C-terminal hydrolase UCH37 (UCH37), mRNA
Homo sapiens angiopoietin-like 4 (ANGPTL4), mRNA
Homo sapiens myotubularin related protein 2 (MTMR2), mRNA
Homo sapiens progesterone receptor membrane component 1 (PGRMC1),
mRNA
Homo sapiens nuclear receptor co-repressor 2 (NCOR2), mRNA
Homo sapiens progesterone receptor membrane component 2 (PGRMC2),
mRNA
Homo sapiens solute carrier family 26, member 4 (SLC26A4), mRNA
Homo sapiens Rho guanine nucleotide exchange factor (GEF) 4 (ARHGEF4),
transcript variant 2, mRNA
Homo sapiens Rho guanine nucleotide exchange factor (GEF) 4 (ARHGEF4), transcript variant 1, mRNA
Homo sapiens Rho guanine exchange factor (GEF) 16 (ARHGEF16), mRNA
Homo sapiens Rho guanine nucleotide exchange factor (GEF) 5 (ARHGEF5),
mRNA
Homo sapiens rho/rac guanine nucleotide exchange factor (GEF) 2 (ARHGEF2),
mRNA
Homo sapiens Rho guanine nucleotide exchange factor (GEF) 1 (ARHGEF1),
mRNA
Homo sapiens ribosomal protein S28 (RPS28), mRNA
Homo sapiens ribosomal protein S27 (metallopanstimulin 1) (RPS27), mRNA
Homo sapiens ribosomal protein S26 (RPS26), mRNA
Homo sapiens replication factor C (activator 1) 1 (145kD) (RFC1), mRNA
Homo sapiens GTF2I repeat domain-containing 1 (GTF2IRD1), transcript variant 2, mRNA
Homo sapiens fibroblast growth factor 19 (FGF19), mRNA
Homo sapiens dyskeratosis congenita 1, dyskerin (DKC1), mRNA
Homo sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump)
membrane sector associated protein M8-9 (APT6M8-9), mRNA
Homo sapiens collagen, type VI, alpha 1 (COL6A1), mRNA
Homo sapiens cadherin 6, type 2, K-cadherin (fetal kidney) (CDH6), mRNA
Homo sapiens solute carrier family 25 (mitochondrial carrier; Graves disease
autoantigen), member 16 (SLC25A16), nuclear gene encoding mitochondrial
protein, mRNA
Homò sapiens synaptotagmin-like 2 (SYTL2), transcript variant a, mRNA
Homo sapiens smoothelin (SMTN), mRNA
Homo sapiens holocarboxylase synthetase (biotin-[proprionyl-Coenzyme A-
carboxylase (ATP-hydrolysing)] ligase) (HLCS), mRNA
Homo sapiens solute carrier family 2 (facilitated glucose transporter), member 10
(SLC2A10), mRNA
Homo sapiens RAN binding protein 17 (RANBP17), mRNA
Homo sapiens activity-dependent neuroprotector (ADNP), mRNA
Homo sapiens RAN binding protein 16 (RANBP16), mRNA
Homo sapiens kallikrein 14 (KLK14), mRNA
Homo sapiens solute carrier family 2 (facilitated glucose transporter), member 9
(SLC2A9), mRNA
Homo sapiens fibroblast growth factor 20 (FGF20), mRNA
Homo sapiens Rho guanine nucleotide exchange factor (GEF) 3 (ARHGEF3),
mRNA
Homo sapiens RAB23, member RAS oncogene family (RAB23), mRNA
Homo sapiens Rho guanine nucleotide exchange factor (GEF) 10 (ARHGEF10),

	mRNA
NM 006989	Homo sapiens Ca2+-promoted Ras inactivator (CAPRI), mRNA
NM 006568	Homo sapiens cell growth regulatory with ring finger domain (CGR19), mRNA
NM 004841	Homo sapiens RAS protein activator like 2 (RASAL2), mRNA
NM 004115	Homo sapiens fibroblast growth factor 14 (FGF14), mRNA
NM 003244	Homo sapiens TGFB-induced factor (TALE family homeobox) (TGIF), mRNA
NM 007285	Homo sapiens GABA(A) receptor-associated protein-like 2 (GABARAPL2),
11112_007.200	mRNA
NM_006047	Homo sapiens RNA binding motif protein 12 (RBM12), mRNA
NM 032588	Homo sapiens ring finger protein 28 (RNF28), mRNA
NM 030766	Homo sapiens apoptosis regulator BCL-G (BCLG), mRNA
NM 022788	Homo sapiens Purinergic receptor P2Y, G protein-coupled, 12 (P2RY12),
	mRNA
NM_015641	Homo sapiens testis derived transcript (3 LIM domains) (TES), mRNA
NM 018144	Homo sapiens Sec61 alpha form 2 (FLJ10578), mRNA
NM_032015	Homo sapiens ring finger protein 26 (RNF26), mRNA
NM_014713	Homo sapiens lysosomal-associated protein transmembrane 4 alpha
_	(LAPTM4A), mRNA
NM_020415	Homo sapiens found in inflammatory zone 3 (FIZZ3), mRNA
NM_020358	Homo sapiens ring finger protein 18 (RNF18), mRNA
NM_005882	Homo sapiens macrophage erythroblast attacher (MAEA), mRNA
NM_016523	Homo sapiens killer cell lectin-like receptor subfamily F, member 1 (KLRF1),
	mRNA
NM_014141	Homo sapiens contactin associated protein-like 2 (CNTNAP2), mRNA
NM_006862	Homo sapiens tudor and KH domain-containing protein (TDRKH), mRNA
NM_006779	Homo sapiens Cdc42 effector protein 2 (CEP2), mRNA
NM_006292	Homo sapiens tumor susceptibility gene 101 (TSG101), mRNA
NM_006449	Homo sapiens Cdc42 effector protein 3 (CEP3), mRNA
NM_002558	Homo sapiens purinergic receptor P2X, ligand-gated ion channel, 1 (P2RX1), mRNA
NM 006712	Homo sapiens FAST kinase (FASTK), transcript variant 1, mRNA
NM 033015	Homo sapiens FAST kinase (FASTK), transcript variant 2, mRNA
NM 025096	Homo sapiens FAST kinase (FASTK), transcript variant 3, mRNA
NM 003852	Homo sapiens transcriptional intermediary factor 1 (TIF1), mRNA
NM 003770	Homo sapiens keratin, hair, acidic, 7 (KRTHA7), mRNA
NM 021013	Homo sapiens keratin, hair, acidic, 4 (KRTHA4), mRNA
NM_004068	Homo sapiens adaptor-related protein complex 2, mu 1 subunit (AP2M1), mRNA
NM_006803	Homo sapiens adaptor-related protein complex 3, mu 2 subunit (AP3M2), mRNA
NM_005498	Homo sapiens adaptor-related protein complex 1, mu 2 subunit (AP1M2), mRNA
NM 032981	Homo sapiens dystrobrevin, alpha (DTNA), transcript variant zeta, mRNA
NM 032980	Homo sapiens dystrobrevin, alpha (DTNA), transcript variant epsilon, mRNA
NM 032979	Homo sapiens dystrobrevin, alpha (DTNA), transcript variant gamma, mRNA
NM 032978	Homo sapiens dystrobrevin, alpha (DTNA), transcript variant beta, mRNA
NM 032975	Homo sapiens dystrobrevin, alpha (DTNA), transcript variant alpha, mRNA
NM 001392	Homo sapiens dystrobrevin, alpha (DTNA), transcript variant DTN3, mRNA
NM 001391	Homo sapiens dystrobrevin, alpha (DTNA), transcript variant DTN2, mRNA
NM 001390	Homo sapiens dystrobrevin, alpha (DTNA), transcript variant DTN1, mRNA
NM 001026	Homo sapiens ribosomal protein S24 (RPS24), transcript variant 2, mRNA
NM 033022	Homo sapiens ribosomal protein S24 (RPS24), transcript variant 1, mRNA
1111 055022	1 120110 captons 11000011111 protein 52 . (14 024), transcript variant 1, mid 14

	·
NM_024416	Homo sapiens osteoglycin (osteoinductive factor, mimecan) (OGN), transcript variant 2, mRNA
NM_033014	Homo sapiens osteoglycin (osteoinductive factor, mimecan) (OGN), transcript variant 1, mRNA
NM_014057	Homo sapiens osteoglycin (osteoinductive factor, mimecan) (OGN), transcript variant 3, mRNA
NM_016152	Homo sapiens retinoic acid receptor, beta (RARB), transcript variant 2, mRNA
NM_000965	Homo sapiens retinoic acid receptor, beta (RARB), transcript variant 1, mRNA
NM_032977	Homo sapiens caspase 10, apoptosis-related cysteine protease (CASP10), transcript variant D, mRNA
NM_032976	Homo sapiens caspase 10, apoptosis-related cysteine protease (CASP10), transcript variant C, mRNA
NM_032974	Homo sapiens caspase 10, apoptosis-related cysteine protease (CASP10), transcript variant B, mRNA
NM_001230	Homo sapiens caspase 10, apoptosis-related cysteine protease (CASP10), transcript variant A, mRNA
NM_032992	Homo sapiens caspase 6, apoptosis-related cysteine protease (CASP6), transcript variant beta, mRNA
NM_001226	Homo sapiens caspase 6, apoptosis-related cysteine protease (CASP6), transcript variant alpha, mRNA
NM_033133	Homo sapiens 2',3'-cyclic nucleotide 3' phosphodiesterase (CNP), mRNA
NM 033125	Homo sapiens organic cation transporter OKB1 (OKB1), mRNA
NM_020349	Homo sapiens ankyrin repeat domain 2 (stretch responsive muscle) (ANKRD2), mRNA
NM_000540	Homo sapiens ryanodine receptor 1 (skeletal) (RYR1), mRNA
NM_016930	Homo sapiens syntaxin 18 (STX18), mRNA
NM_014808	Homo sapiens KIAA0793 gene product (KIAA0793), mRNA
NM_005428	Homo sapiens vav 1 oncogene (VAV1), mRNA
NM_005747	Homo sapiens elastase 3A, pancreatic (protease E) (ELA3A), mRNA
NM_000922	Homo sapiens phosphodiesterase 3B, cGMP-inhibited (PDE3B), mRNA
NM_033069	Homo sapiens ADG-90 protein (ADG-90), mRNA
NM_033085	Homo sapiens fetal and adult testis expressed transcript protein (FATE), mRNA
NM_015001	Homo sapiens SMART/HDAC1 associated repressor protein (SHARP), mRNA
NM_032984	Homo sapiens caspase 2, apoptosis-related cysteine protease (neural precursor cell expressed, developmentally down-regulated 2) (CASP2), transcript variant 4, mRNA
NM_032983	Homo sapiens caspase 2, apoptosis-related cysteine protease (neural precursor cell expressed, developmentally down-regulated 2) (CASP2), transcript variant 3, mRNA
NM_032982	Homo sapiens caspase 2, apoptosis-related cysteine protease (neural precursor cell expressed, developmentally down-regulated 2) (CASP2), transcript variant 1, mRNA
NM_032957	Homo sapiens tumor necrosis factor receptor superfamily, member 6b, decoy (TNFRSF6B), transcript variant 1, mRNA
NM_032945	Homo sapiens tumor necrosis factor receptor superfamily, member 6b, decoy (TNFRSF6B), transcript variant M68C, mRNA
NM_001224	Homo sapiens caspase 2, apoptosis-related cysteine protease (neural precursor cell expressed, developmentally down-regulated 2) (CASP2), transcript variant 2, mRNA
NM_015647	Homo sapiens tumor necrosis factor receptor superfamily, member 6b, decoy (TNFRSF6B), transcript variant 3, mRNA
NM_033012	Homo sapiens tumor necrosis factor (ligand) superfamily, member 11

	L (TD) TD(CF) (A)
) D. C. 000701	(TNFSF11), transcript variant 2, mRNA
NM_003701	Homo sapiens tumor necrosis factor (ligand) superfamily, member 11
27.5 005.100	(TNFSF11), transcript variant 1, mRNA
NM_005409	Homo sapiens small inducible cytokine subfamily B (Cys-X-Cys), member 11 (SCYB11), mRNA
NM_005035	Homo sapiens polymerase (RNA) mitochondrial (DNA directed) (POLRMT), nuclear gene encoding mitochondrial protein, mRNA
NM_006980	Homo sapiens transcription termination factor, mitochondrial (MTERF), nuclear
	gene encoding mitochondrial protein, mRNA
NM_001305	Homo sapiens claudin 4 (CLDN4), mRNA
NM_032996	Homo sapiens caspase 9, apoptosis-related cysteine protease (CASP9), transcript variant beta, mRNA
NM_001229	Homo sapiens caspase 9, apoptosis-related cysteine protease (CASP9), transcript variant alpha, mRNA
NM 004346	Homo sapiens caspase 3, apoptosis-related cysteine protease (CASP3), transcript
_	variant alpha, mRNA
NM_032991	Homo sapiens caspase 3, apoptosis-related cysteine protease (CASP3), transcript variant beta, mRNA
NM_033057	Homo sapiens olfactory receptor, family 2, subfamily B, member 2 (OR2B2),
	mRNA
NM_033051	Homo sapiens thymic stromal co-transporter (TSCOT), mRNA
NM_033048	Homo sapiens CPX chromosome region, candidate 1 (CPXCR1), mRNA
NM_033007	Homo sapiens death effector filament-forming Ced-4-like apoptosis protein (DEFCAP), transcript variant E, mRNA
NM 033006	Homo sapiens death effector filament-forming Ced-4-like apoptosis protein
	(DEFCAP), transcript variant D, mRNA
NM_033005	Homo sapiens death effector filament-forming Ced-4-like apoptosis protein (DEFCAP), transcript variant C, mRNA
NM 033004	Homo sapiens death effector filament-forming Ced-4-like apoptosis protein
1111_0000,	(DEFCAP), transcript variant A, mRNA
NM_014922	Homo sapiens death effector filament-forming Ced-4-like apoptosis protein
	(DEFCAP), transcript variant B, mRNA
NM 000088	Homo sapiens collagen, type I, alpha 1 (COL1A1), mRNA
NM_019105	Homo sapiens tenascin XB (TNXB), transcript variant XB, mRNA
NM 033036	Homo sapiens beta-galactose-3-O-sulfotransferase 3 (GAL3ST2), mRNA
NM_033029	Homo sapiens leishmanolysin-like (metallopeptidase M8 family) (LMLN),
11111_00000	mRNA
NM 033028	Homo sapiens Bardet-Biedl syndrome 4 (BBS4), mRNA
NM 021807	Homo sapiens secretory protein SEC8 (SEC8), mRNA
NM 020137	Homo sapiens GRIP-associated protein 1 (GRASP1), mRNA
NM 015133	Homo sapiens mitogen-activated protein kinase 8 interacting protein 3
	(MAPK8IP3), mRNA
NM 014006	Homo sapiens PI-3-kinase-related kinase SMG-1 (SMG1), mRNA
NM_021914	Homo sapiens cofilin 2 (muscle) (CFL2), mRNA
NM 032520	Homo sapiens hypothetical protein CAB56184 (CAB56184), mRNA
NM 032923	Homo sapiens hypothetical protein MGC16025 (MGC16025), mRNA
NM_032917	Homo sapiens hypothetical protein MGC2848 (MGC2848), mRNA
NM 032868	Homo sapiens hypothetical protein FLJ14981 (FLJ14981), mRNA
NM 032862	Homo sapiens hypothetical protein FLJ14926 (FLJ14926), mRNA
NM 032801	Homo sapiens hypothetical protein FLJ14529 (FLJ14529), mRNA
NM 032753	Homo sapiens hypothetical protein MGC15631 (MGC15631), mRNA
	Homo sapiens hypothetical protein MGC2721 (MGC2721) mDNA
NM_032753 NM_032737	Homo sapiens hypothetical protein MGC15631 (MGC15631), mRNA Homo sapiens hypothetical protein MGC2721 (MGC2721), mRNA

NM_032668	Homo sapiens hypothetical protein MGC4771 (MGC4771), mRNA
NM_032503	Homo sapiens G protein-coupled receptor slt (SLT), mRNA
NM_032377	Homo sapiens hypothetical protein MGC4549 (MGC4549), mRNA
NM_032326	Homo sapiens hypothetical protein MGC4618 (MGC4618), mRNA
NM 032306	Homo sapiens hypothetical protein MGC10974 (MGC10974), mRNA
NM 032281	Homo sapiens hypothetical protein DKFZp547J036 (DKFZp547J036), mRNA
NM 015650	Homo sapiens microtubule-interacting protein that associates with TRAF3 (MIP-
_	T3), mRNA
NM_031487	Homo sapiens hypothetical protein MGC4604 (MGC4604), mRNA
NM_031470	Homo sapiens junctional adhesion molecule 3 (JAM3), mRNA
NM_031304	Homo sapiens hypothetical protein MGC4293 (MGC4293), mRNA
NM 031213	Homo sapiens hypothetical protein MGC:5244, (MGC:5244), mRNA
NM 031208	Homo sapiens hypothetical protein DKFZp566J2046 (DKFZP566J2046), mRNA
NM 030924	Homo sapiens hypothetical protein PRTD-NY3 (PRTD-NY3), mRNA
NM 030824	Homo sapiens hypothetical protein FLJ14356 (FLJ14356), mRNA
NM 030631	Homo sapiens solute carrier family 25 (mitochondrial oxodicarboxylate carrier),
, —	member 21 (SLC25A21), mRNA
NM 024571	Homo sapiens hypothetical protein FLJ22940 (FLJ22940), mRNA
NM 025015	Homo sapiens KIAA0417 gene product (KIAA0417), mRNA
NM 024103	Homo sapiens hypothetical protein MGC2615 (MGC2615), mRNA
NM 030578	Homo sapiens hypothetical protein MGC4093 (MGC4093), mRNA
NM 014015	Homo sapiens MYLE protein (MYLE), mRNA
NM 025094	Homo sapiens hypothetical protein FLJ22184 (FLJ22184), mRNA
NM 025078	Homo sapiens hypothetical protein FLJ22378 (FLJ22378), mRNA
NM 025061	Homo sapiens hypothetical protein FLJ23420 (FLJ23420), mRNA
NM 024967	Homo sapiens hypothetical protein FLJ11637 (FLJ11637), mRNA
NM 024898	Homo sapiens hypothetical protein FLJ22757 (FLJ22757), mRNA
NM 024877	Homo sapiens hypothetical protein FLJ13265 (FLJ13265), mRNA
NM 024726	Homo sapiens hypothetical protein FLJ22527 (FLJ22527), mRNA
NM 024719	Homo sapiens hypothetical protein FLJ22474 (FLJ22474), mRNA
NM_024600	Homo sapiens hypothetical protein FLJ20898 (FLJ20898), mRNA
NM_024508	Homo sapiens hypothetical protein MGC10796 (MGC10796), mRNA
NM 024341	Homo sapiens hypothetical protein MGC4054 (MGC4054), mRNA
NM_024064	Homo sapiens hypothetical protein MGC5363 (MGC5363), mRNA
NM_024029	Homo sapiens hypothetical protein MGC3262 (MGC3262), mRNA
NM_023078	Homo sapiens hypothetical protein FLJ13852 (FLJ13852), mRNA
NM_023076	Homo sapiens hypothetical protein FLJ23360 (FLJ23360), mRNA
NM_022842	Homo sapiens hypothetical protein FLJ22969 (FLJ22969), mRNA
NM_022737	Homo sapiens hypothetical protein FLJ13055 (FLJ13055), mRNA
NM 022459	Homo sapiens hypothetical protein FLJ13046 similar to exportin 4; KIAA1721
_	pr (FLJ13046), mRNA
NM 022437	Homo sapiens ATP-binding cassette, sub-family G (WHITE), member 8 (sterolin
_	2) (ABCG8), mRNA
NM 022135	Homo sapiens popeye protein 2 (POP2), mRNA
NM 022066	Homo sapiens likely ortholog of mouse ubiquitin-conjugating enzyme E2-230K
_	(E2-230K), mRNA
NM_015480	Homo sapiens nectin 3 (DKFZP566B0846), mRNA
NM 004240	Homo sapiens thyroid hormone receptor interactor 10 (TRIP10), mRNA
NM_003589	Homo sapiens cullin 4A (CUL4A), mRNA
NM_021731	Homo sapiens hypothetical protein PP3501 (PP3501), mRNA
NM 020129	Homo sapiens placental protein 13-like protein (LOC56891), mRNA
NM 020196	Homo sapiens HCNP protein; XPA-binding protein 2 (HCNP), mRNA
	1 deploted House, 22 oman's protein 2 (HOIL), HIGH

NM_020224	Homo sapiens hypothetical protein DKFZp547O146 (DKFZp547O146), mRNA
NM_019064	Homo sapiens hypothetical protein (FLJ10832), mRNA
NM_019012	Homo sapiens phosphoinositol 3-phosphate-binding protein-2 (PEPP2), mRNA
NM_018635	Homo sapiens hypothetical protein PRO2900 (PRO2900), mRNA
NM_018687	Homo sapiens hepatocellular carcinoma-associated gene TD26 (LOC55908), mRNA
NM_018441	Homo sapiens peroxisomal trans 2-enoyl CoA reductase; putative short chain alcohol dehydrogenase (HSA250303), mRNA
NM 018645	Homo sapiens hypothetical protein HES6 (HES6), mRNA
NM 017967	Homo sapiens hypothetical protein FLJ20850 (FLJ20850), mRNA
NM 017914	Homo sapiens hypothetical protein FLJ20640 (FLJ20640), mRNA
NM 017905	Homo sapiens hypothetical protein FLJ20623 (FLJ20623), mRNA
NM 017722	Homo sapiens hypothetical protein FLJ20244 (FLJ20244), mRNA
NM 017668	Homo sapiens LIS1-interacting protein NUDE1, rat homolog (NUDE1), mRNA
NM 017616	Homo sapiens hypothetical protein FLJ20004 (FLJ20004), mRNA
NM 018185	Homo sapiens hypothetical protein FLJ10704 (FLJ10704), mRNA
NM 018074	Homo sapiens hypothetical protein FLJ10374 (FLJ10374), mRNA
NM 018057	Homo sapiens homolog of rat orphan transporter v7-3 (NTT73), mRNA
NM 018049	Homo sapiens hypothetical protein FLJ10297 (FLJ10297), mRNA
NM 018028	Homo sapiens hypothetical protein FLJ10211 (FLJ10211), mRNA
NM 018000	Homo sapiens hypothetical protein FLJ10116 (FLJ10116), mRNA
NM 016510	Homo sapiens putative selenocysteine lyase (SCLY), mRNA
NM 016434	Homo sapiens tumor necrosis factor receptor superfamily, member 6b, decoy
	(TNFRSF6B), transcript variant 2, mRNA
NM_016289	Homo sapiens MO25 protein (LOC51719), mRNA
NM_016264	Homo sapiens GIOT-2 for gonadotropin inducible transcription repressor-2 (GIOT-2), mRNA
NM_016149	Homo sapiens protein inhibitor of activated STAT protein PIASy (PIASY), mRNA
NM_015897	Homo sapiens protein inhibitor of activated STAT protein PIASy (PIASY), mRNA
NM 016581	Homo sapiens ECSIT (LOC51295), mRNA
NM 016479	Homo sapiens hypothetical protein (LOC51246), mRNA
NM 016474	Homo sapiens hypothetical protein (LOC51244), mRNA
NM 016094	Homo sapiens HSPC042 protein (LOC51122), mRNA
NM 015942	Homo sapiens CGI-12 protein (LOC51001), mRNA
NM 016475	Homo sapiens hypothetical protein (HSPC213), mRNA
NM 016457	Homo sapiens protein kinase D2 (PKD2), mRNA
NM 016111	Homo sapiens KIAA0683 gene product (KIAA0683), mRNA
NM 014049	Homo sapiens NPD002 protein (NPD002), mRNA
NM 014963	Homo sapiens KIAA0963 protein (KIAA0963), mRNA
NM 015571	Homo sapiens SUMO-1-specific protease (SUSP1), mRNA
NM 014789	Homo sapiens KIAA0628 gene product (KIAA0628), mRNA
NM 014714	Homo sapiens KIAA0590 gene product (KIAA0590), mRNA
NM 014714	Homo sapiens KIAA0254 gene product (KIAA0254), mRNA
NM 014065	Homo sapiens HT001 protein (HT001), mRNA
NM 014170	Homo sapiens HSPC135 protein (HSPC135), mRNA
NM 015462	Homo sapiens DKFZP586L0724 protein (DKFZP586L0724), mRNA
	Homo sapiens zinc finger protein 288 (ZNF288), mRNA
NM_015642	Homo sapiens DKFZP434N161 protein (DKFZP434N161), mRNA
NM_015493	Homo sapiens DKFZF434N161 protein (DKFZF434N161), mKNA  Homo sapiens muscle-specific beta 1 integrin binding protein (MIBP), mRNA
NM_014446	
NM_013314	Homo sapiens B-cell linker (BLNK), mRNA

NM_007086	Homo sapiens AND-1 protein (AND-1), mRNA
NM_006701	Homo sapiens similar to S. pombe dim1+ (DIM1), mRNA
NM_006300	Homo sapiens zinc finger protein 230 (ZNF230), mRNA
NM_006477	Homo sapiens RAS-related on chromosome 22 (RRP22), mRNA
NM_006087	Homo sapiens tubulin, beta, 5 (TUBB5), mRNA
NM_006056	Homo sapiens G protein-coupled receptor 66 (GPR66), mRNA
NM_005815	Homo sapiens Kruppel-type zinc finger (C2H2) (ZK1), mRNA
NM_005817	Homo sapiens cargo selection protein (mannose 6 phosphate receptor binding protein) (TIP47), mRNA
NM_005801	Homo sapiens putative translation initiation factor (SUI1), mRNA
NM_005837	Homo sapiens POP7 (processing of precursor, S. cerevisiae) homolog (RPP20), mRNA
NM_005776	Homo sapiens cornichon-like (CNIL), mRNA
NM_004970	Homo sapiens insulin-like growth factor binding protein, acid labile subunit (IGFALS), mRNA
NM_004945	Homo sapiens dynamin 2 (DNM2), mRNA
NM_004283	Homo sapiens RAB3D, member RAS oncogene family (RAB3D), mRNA
NM_004548	Homo sapiens NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 10 (22kD, PDSW) (NDUFB10), mRNA
NM 004124	Homo sapiens glia maturation factor, beta (GMFB), mRNA
NM 004877	Homo sapiens glia maturation factor, gamma (GMFG), mRNA
NM 004907	Homo sapiens immediate early protein (ETR101), mRNA
NM 004044	Homo sapiens 5-aminoimidazole-4-carboxamide ribonucleotide
_	formyltransferase/IMP cyclohydrolase (ATIC), mRNA
NM_004315	Homo sapiens N-acylsphingosine amidohydrolase (acid ceramidase) (ASAH), mRNA
NM_004846	Homo sapiens eukaryotic translation initiation factor 4E-like 3 (EIF4EL3), mRNA
NM_003765	Homo sapiens syntaxin 10 (STX10), mRNA
NM_003110	Homo sapiens Sp2 transcription factor (SP2), mRNA
NM_003113	Homo sapiens nuclear antigen Sp100 (SP100), mRNA
NM_000543	Homo sapiens sphingomyelin phosphodiesterase 1, acid lysosomal (acid sphingomyelinase) (SMPD1), mRNA
NM_003072	Homo sapiens SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 4 (SMARCA4), mRNA
NM_002807	Homo sapiens proteasome (prosome, macropain) 26S subunit, non-ATPase, 1 (PSMD1), mRNA
NM_002704	Homo sapiens pro-platelet basic protein (includes platelet basic protein, beta-thromboglobulin, connective tissue-activating peptide III, neutrophil-activating peptide-2) (PPBP), mRNA
NM_000089	Homo sapiens collagen, type I, alpha 2 (COL1A2), mRNA
NM_001687	Homo sapiens ATP synthase, H+ transporting, mitochondrial F1 complex, delta subunit (ATP5D), mRNA
NM_020168	Homo sapiens p21(CDKN1A)-activated kinase 6 (PAK6), mRNA
NM_032657	Homo sapiens hypothetical protein MGC10442 (MGC10442), mRNA
NM_032571	Homo sapiens EGF-like module-containing mucin-like receptor EMR3 (EMR3), mRNA
NM_032413	Homo sapiens normal mucosa of esophagus specific 1 (NMES1), mRNA
NM_015093	Homo sapiens TAK1-binding protein 2 (TAB2), mRNA
NM_031947	Homo sapiens ornithine transporter 2 (ORNT2), mRNA
	- Control of Control o
NM_005563	Homo sapiens stathmin 1/oncoprotein 18 (STMN1) mRNA
NM_005563 NM_024662	Homo sapiens stathmin 1/oncoprotein 18 (STMN1), mRNA Homo sapiens hypothetical protein FLJ10774 (FLJ10774), mRNA

NM_024637	Homo sapiens beta-galactose-3-O-sulfotransferase, 4 (GAL3ST-4), mRNA
NM 024617	Homo sapiens hypothetical protein FLJ13409 (FLJ13409), mRNA
NM 020796	Homo sapiens sema domain, transmembrane domain (TM), and cytoplasmic
	domain, (semaphorin) 6A (SEMA6A), mRNA
NM 013283	Homo sapiens methionine adenosyltransferase II, beta (MAT2B), mRNA
NM 012231	Homo sapiens PR domain containing 2, with ZNF domain (PRDM2), mRNA
NM 020428	Homo sapiens CTL2 gene (CTL2), mRNA
NM 015866	Homo sapiens PR domain containing 2, with ZNF domain (PRDM2), mRNA
NM 014771	Homo sapiens 95 kDa retinoblastoma protein binding protein; KIAA0661 gene
_	pro (KIAA0661), mRNA
NM 014454	Homo sapiens p53 regulated PA26 nuclear protein (PA26), mRNA
NM 013447	Homo sapiens egf-like module containing, mucin-like, hormone receptor-like
	sequence 2 (EMR2), mRNA
NM_006499	Homo sapiens lectin, galactoside-binding, soluble, 8 (galectin 8) (LGALS8),
	mRNA
NM 006031	Homo sapiens pericentrin 2 (kendrin) (PCNT2), mRNA
NM 022040	Homo sapiens Williams-Beuren syndrome chromosome region 5 (WBSCR5),
1414_022010	transcript variant 1, mRNA
NM_032464	Homo sapiens Williams-Beuren syndrome chromosome region 5 (WBSCR5),
1441_032-10-1	transcript variant 4, mRNA
NM_032463	Homo sapiens Williams-Beuren syndrome chromosome region 5 (WBSCR5),
1111_032103	transcript variant 2, mRNA
NM_014146	Homo sapiens Williams-Beuren syndrome chromosome region 5 (WBSCR5),
1111_01 11 10	transcript variant 3, mRNA
NM_031992	Homo sapiens Williams-Beuren syndrome chromosome region 1 (WBSCR1),
1111_051552	transcript variant 2, mRNA
NM 006234	Homo sapiens polymerase (RNA) II (DNA directed) polypeptide J (13.3kD)
11112_000220,	(POLR2J), transcript variant a, mRNA
NM_032959	Homo sapiens polymerase (RNA) II (DNA directed) polypeptide J (13.3kD)
11212_00200	(POLR2J), transcript variant b, mRNA
NM_032958	Homo sapiens polymerase (RNA) II (DNA directed) polypeptide J (13.3kD)
	(POLR2J), transcript variant c, mRNA
NM 002694	Homo sapiens polymerase (RNA) II (DNA directed) polypeptide C (33kD)
_	(POLR2C), transcript variant alpha, mRNA
NM 032940	Homo sapiens polymerase (RNA) II (DNA directed) polypeptide C (33kD)
	(POLR2C), transcript variant gamma, mRNA
NM_033011	Homo sapiens plasminogen activator, tissue (PLAT), transcript variant 3, mRNA
NM 000931	Homo sapiens plasminogen activator, tissue (PLAT), transcript variant 2, mRNA
NM 000930	Homo sapiens plasminogen activator, tissue (PLAT), transcript variant 1, mRNA
NM 033013	Homo sapiens nuclear receptor subfamily 1, group I, member 2 (NR1I2),
	transcript variant 3, mRNA
NM 003889	Homo sapiens nuclear receptor subfamily 1, group I, member 2 (NR1I2),
_	transcript variant 1, mRNA
NM 022002	Homo sapiens nuclear receptor subfamily 1, group I, member 2 (NR1I2),
	transcript variant 2, mRNA
NM 022170	Homo sapiens Williams-Beuren syndrome chromosome region 1 (WBSCR1),
	transcript variant 1, mRNA
NM 032408	Homo sapiens bromodomain adjacent to zinc finger domain, 1B (BAZ1B),
	transcript variant 2, mRNA
NM 023005	Homo sapiens bromodomain adjacent to zinc finger domain, 1B (BAZ1B),
	transcript variant 1, mRNA
NM 001024	Homo sapiens ribosomal protein S21 (RPS21), mRNA
	1 Captolio Hoodellias provincias (24 551), sint 111

tagonizing transcription factor (DED), mRNA
protein F (350/400kD, mitosin) (CENPF), mRNA
peta)-like 2 (TBL2), transcript variant 2, mRNA
rotein 278 (ZNF278), transcript variant 3, mRNA
rotein 278 (ZNF278), transcript variant 4, mRNA
rotein 278 (ZNF278), transcript variant 2, mRNA
rotein 278 (ZNF278), transcript variant 1, mRNA
scription factor II, i (GTF2I), transcript variant 5,
, , , , , , , , , , , , , , , , , , , ,
scription factor II, i (GTF2I), transcript variant 4,
scription factor II, i (GTF2I), transcript variant 3,
scription factor II, i (GTF2I), transcript variant 2,
scription factor II, i (GTF2I), transcript variant 1,
nding protein (RDBP), mRNA
ivated protein kinase kinase 1 (MAP2K1), mRNA
beta)-like 2 (TBL2), transcript variant 1, mRNA
lyl isomerase H (cyclophilin H) (PPIH), mRNA
sphatase, intestinal (ALPI), mRNA
octanoyltransferase (CROT), mRNA
se A2, group IVB (cytosolic) (PLA2G4B), mRNA
air cross-complementing rodent repair deficiency,
ERCC6), mRNA
TOR), mRNA
l (PB1), mRNA
l (PB1), mRNA
al ribosomal protein L30 (MRPL30), mRNA
sus associated putative guanine nucleotide exchange f
ituent protein (MSE55), mRNA
MYO1A), mRNA
P450, subfamily XXIA (steroid 21-hydroxylase,
asia), polypeptide 2 (CYP21A2), mRNA
component 2 (C2), mRNA
al ribosomal protein L13 (MRPL13), mRNA
al ribosomal protein L23 (MRPL23), mRNA
n-like and metalloprotease (reprolysin type) with
tif, 9 (ADAMTS9), mRNA
se transition 2 (GSPT2), mRNA
al ribosomal protein L22 (MRPL22), mRNA
al ribosomal protein L15 (MRPL15), mRNA
n protein 5 (ponsin) (SH3D5), mRNA
n protein 5 (ponsin) (SH3D5), mRNA
emia, complementation group A (FANCA), mRNA
ane protease, serine 2 (TMPRSS2), mRNA
(RNA) II (DNA directed) polypeptide F (POLR2F),
, , , , , , , , , , , , , , , , , , , ,
ible cytokine subfamily A (Cys-Cys), member 15

NM_032965	Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 15 (SCYA15), transcript variant 3, mRNA
NM 032964	Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 15
	(SCYA15), transcript variant 1, mRNA
NM_032454	Homo sapiens serine/threonine kinase 19 (STK19), transcript variant 2, mRNA
NM_007057	Homo sapiens ZW10 interactor (ZWINT), transcript variant 1, mRNA
NM_032997	Homo sapiens ZW10 interactor (ZWINT), transcript variant 2, mRNA
NM_003262	Homo sapiens translocation protein 1 (TLOC1), mRNA
NM_032470	Homo sapiens tenascin XB (TNXB), transcript variant XB-S, mRNA
NM_004166	Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 14
	(SCYA14), transcript variant 1, mRNA
NM_032963	Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 14
	(SCYA14), transcript variant 3, mRNA
NM_032962	Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 14
	(SCYA14), transcript variant 2, mRNA
NM_021219	Homo sapiens junctional adhesion molecule 2 (JAM2), mRNA
NM_014456	Homo sapiens programmed cell death 4 (neoplastic transformation inhibitor)
	(PDCD4), mRNA
NM_004197	Homo sapiens serine/threonine kinase 19 (STK19), transcript variant 1, mRNA
NM_007214	Homo sapiens SEC63, endoplasmic reticulum translocon component (S.
	cerevisiae (SEC63L), mRNA
NM_006808	Homo sapiens protein translocation complex beta (SEC61B), mRNA
NM_001028	Homo sapiens ribosomal protein S25 (RPS25), mRNA
NM_001022	Homo sapiens ribosomal protein S19 (RPS19), mRNA
NM_001021	Homo sapiens ribosomal protein S17 (RPS17), mRNA
NM_001020	Homo sapiens ribosomal protein S16 (RPS16), mRNA
NM_001018	Homo sapiens ribosomal protein S15 (RPS15), mRNA
NM_001017	Homo sapiens ribosomal protein S13 (RPS13), mRNA
NM_012423	Homo sapiens ribosomal protein L13a (RPL13A), mRNA
NM_002907	Homo sapiens RecQ protein-like (DNA helicase Q1-like) (RECQL), transcript variant 1, mRNA
NM_032941	Homo sapiens RecQ protein-like (DNA helicase Q1-like) (RECQL), transcript
_	variant 2, mRNA
NM_021128	Homo sapiens polymerase (RNA) II (DNA directed) polypeptide L (7.6kD) (POLR2L), mRNA
NM_006233	Homo sapiens polymerase (RNA) II (DNA directed) polypeptide I (14.5kD)
	(POLR2I), mRNA
NM_006232	Homo sapiens polymerase (RNA) II (DNA directed) polypeptide H (POLR2H),
_	mRNA
NM_002695	Homo sapiens polymerase (RNA) II (DNA directed) polypeptide E (25kD)
	(POLR2E), mRNA
NM_004805	Homo sapiens polymerase (RNA) II (DNA directed) polypeptide D (POLR2D),
	mRNA
NM_000937	Homo sapiens polymerase (RNA) II (DNA directed) polypeptide A (220kD)
	(POLR2A), mRNA
NM_001987	Homo sapiens ets variant gene 6 (TEL oncogene) (ETV6), mRNA
NM_032973	Homo sapiens protocadherin 22 (PCDH22), transcript variant c, mRNA
NM_032972	Homo sapiens protocadherin 22 (PCDH22), transcript variant b, mRNA
NM_032971	Homo sapiens protocadherin 22 (PCDH22), transcript variant a, mRNA
NM_020403	Homo sapiens protocadherin 9 (PCDH9), mRNA
NM_022843	Homo sapiens protocadherin 20 (PCDH20), mRNA
NM_032949	Homo sapiens protocadherin 8 (PCDH8), transcript variant 2, mRNA
	, may 2,

NM_032457	Homo sapiens BH-protocadherin (brain-heart) (PCDH7), transcript variant c, mRNA
NM_032456	Homo sapiens BH-protocadherin (brain-heart) (PCDH7), transcript variant b, mRNA
NM_002589	Homo sapiens BH-protocadherin (brain-heart) (PCDH7), transcript variant a, mRNA
NM_016580	Homo sapiens protocadherin 12 (PCDH12), mRNA
NM_032420	Homo sapiens protocadherin 1 (cadherin-like 1) (PCDH1), transcript variant 2, mRNA
NM_032969	Homo sapiens protocadherin 11 (PCDH11), transcript variant d, mRNA
NM_032968	Homo sapiens protocadherin 11 (PCDH11), transcript variant c, mRNA
NM_032967	Homo sapiens protocadherin 11 (PCDH11), transcript variant b, mRNA
NM_032950	Homo sapiens matrix metalloproteinase 28 (MMP28), transcript variant 2, mRNA
NM_024302	Homo sapiens matrix metalloproteinase 28 (MMP28), transcript variant 1, mRNA
NM_006575	Homo sapiens mitogen-activated protein kinase kinase kinase kinase 5 (MAP4K5), mRNA
NM_004635	Homo sapiens mitogen-activated protein kinase-activated protein kinase 3 (MAPKAPK3), mRNA
NM_002587	Homo sapiens protocadherin 1 (cadherin-like 1) (PCDH1), transcript variant 1, mRNA
NM_004759	Homo sapiens mitogen-activated protein kinase-activated protein kinase 2 (MAPKAPK2), transcript variant 1, mRNA
NM_032960	Homo sapiens mitogen-activated protein kinase-activated protein kinase 2 (MAPKAPK2), transcript variant 2, mRNA
NM 032515	Homo sapiens Bcl-2-related ovarian killer protein-like (BOKL), mRNA
NM 015166	Homo sapiens KIAA0027 protein (MLC1), mRNA
NM_001795	Homo sapiens cadherin 5, type 2, VE-cadherin (vascular epithelium) (CDH5), mRNA
NM_001794	Homo sapiens cadherin 4, type 1, R-cadherin (retinal) (CDH4), mRNA
NM_001793	Homo sapiens cadherin 3, type 1, P-cadherin (placental) (CDH3), mRNA
NM_001792	Homo sapiens cadherin 2, type 1, N-cadherin (neuronal) (CDH2), mRNA
NM 004360	Homo sapiens cadherin 1, type 1, E-cadherin (epithelial) (CDH1), mRNA
NM 006137	Homo sapiens CD7 antigen (p41) (CD7), mRNA
NM_005864	Homo sapiens signal transduction protein (SH3 containing) (EFS2), transcript variant 1, mRNA
NM_032459	Homo sapiens signal transduction protein (SH3 containing) (EFS2), transcript variant 2, mRNA
NM_032107	Homo sapiens lethal (3) malignant brain tumor l(3)mbt protein (Drosophila) ho (H-L(3)MBT), transcript variant II, mRNA
NM_015478	Homo sapiens lethal (3) malignant brain tumor l(3)mbt protein (Drosophila) ho (H-L(3)MBT), transcript variant I, mRNA
NM 004318	Homo sapiens aspartate beta-hydroxylase (ASPH), transcript variant 1, mRNA
NM 032468	Homo sapiens aspartate beta-hydroxylase (ASPH), transcript variant 2, mRNA
NM 032467	Homo sapiens aspartate beta-hydroxylase (ASPH), transcript variant 4, mRNA
NM 032466	Homo sapiens aspartate beta-hydroxylase (ASPH), transcript variant 4, inclyA
NM 020164	Homo sapiens aspartate beta-hydroxylase (ASPH), transcript variant 5, mRNA
NM_014217	Homo sapiens potassium channel, subfamily K, member 2 (TREK-1) (KCNK2), mRNA
NM_031498	Homo sapiens guanine nucleotide binding protein (G protein), gamma transducing activity polypeptide 2 (GNGT2), mRNA

NM 031311	Homo sapiens carboxypeptidase, vitellogenic-like (CPVL), mRNA
NM 022768	Homo sapiens RNA binding motif protein 15 (RBM15), mRNA
NM 021797	Homo sapiens eosinophil chemotactic cytokine (TSA1902), mRNA
NM 014330	Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 15A
_	(PPP1R15A), mRNA
NM 014522	Homo sapiens protocadherin 11 (PCDH11), transcript variant a, mRNA
NM 003004	Homo sapiens secreted and transmembrane 1 (SECTM1), mRNA
NM_002696	Homo sapiens polymerase (RNA) II (DNA directed) polypeptide G (POLR2G),
	mRNA
NM_000938	Homo sapiens polymerase (RNA) II (DNA directed) polypeptide B (140kD) (POLR2B), mRNA
NM_001372	Homo sapiens dynein, axonemal, heavy polypeptide 9 (DNAH9), transcript variant 2, mRNA
NM_004215	Homo sapiens estrogen receptor binding site associated, antigen, 9 (EBAG9), mRNA
NM_005111	Homo sapiens crystallin, zeta (quinone reductase)-like 1 (CRYZL1), mRNA
NM_004381	Homo sapiens cAMP responsive element binding protein-like 1 (CREBL1), mRNA
NM_000592	Homo sapiens complement component 4B (C4B), mRNA
NM_007293	Homo sapiens complement component 4A (C4A), mRNA
NM_032603	Homo sapiens lysyl oxidase-like 3 (LOXL3), mRNA
NM_023937	Homo sapiens mitochondrial ribosomal protein L34 (MRPL34), mRNA
NM_022567	Homo sapiens nyctalopin (NYX), mRNA
NM_022467	Homo sapiens carbohydrate (N-acetylgalactosamine 4-0) sulfotransferase 8 (CHST8), mRNA
NM_016557	Homo sapiens orphan seven-transmembrane receptor, chemokine related (VSHK1), mRNA
NM_016116	Homo sapiens ankyrin repeat and SOCS box-containing 4 (ASB4), mRNA
NM_016114	Homo sapiens ankyrin repeat and SOCS box-containing 1 (ASB1), mRNA
NM_016115	Homo sapiens ankyrin repeat and SOCS box-containing 3 (ASB3), mRNA
NM_014398	Homo sapiens lysosomal-associated membrane protein 3 (LAMP3), mRNA
NM_014434	Homo sapiens NADPH-dependent FMN and FAD containing oxidoreductase (NR1), mRNA
NM_004860	Homo sapiens fragile X mental retardation, autosomal homolog 2 (FXR2), mRNA
NM_006850	Homo sapiens interleukin 24 (IL24), mRNA
NM_006541	Homo sapiens thioredoxin-like 2 (TXNL2), mRNA
NM_004662	Homo sapiens dynein, axonemal, heavy polypeptide 9 (DNAH9), transcript variant 1, mRNA
NM_000029	Homo sapiens angiotensinogen (serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 8) (AGT), mRNA
NM_004050	Homo sapiens BCL2-like 2 (BCL2L2), mRNA
NM_004049	Homo sapiens BCL2-related protein A1 (BCL2A1), mRNA
NM_001623	Homo sapiens allograft inflammatory factor 1 (AIF1), transcript variant 3, mRNA
NM_032955	Homo sapiens allograft inflammatory factor 1 (AIF1), transcript variant 1, mRNA
NG_000010	Homo sapiens genomic cytochrome P450, subfamily IIA (phenobarbital-inducible) (CYP2A.2@) on chromosome 19
NM_004847	Homo sapiens allograft inflammatory factor 1 (AIF1), transcript variant 2, mRNA
NM_005452	Homo sapiens chromosome 6 open reading frame 11 (C6orf11), mRNA

NM_031282 Homo sapiens immunoglobulin superfamily receptor translocation associ (IRTA1), mRNA  NM_031281 Homo sapiens immunoglobulin superfamily receptor translocation associ (IRTA2), mRNA  NM_000767 Homo sapiens cytochrome P450, subfamily IIB (phenobarbital-inducible polypeptide 6 (CYP2B6), mRNA  NM_020165 Homo sapiens postreplication repair protein hRAD18p (RAD18), mRNA  NM_001710 Homo sapiens B-factor, properdin (BF), mRNA  NM_021800 Homo sapiens J domain containing protein 1 (JDP1), mRNA  NM_020404 Homo sapiens tumor endothelial marker 1 precursor (TEM1), mRNA	
NM_031281 Homo sapiens immunoglobulin superfamily receptor translocation associ (IRTA2), mRNA  NM_000767 Homo sapiens cytochrome P450, subfamily IIB (phenobarbital-inducible polypeptide 6 (CYP2B6), mRNA  NM_020165 Homo sapiens postreplication repair protein hRAD18p (RAD18), mRNA  NM_021800 Homo sapiens B-factor, properdin (BF), mRNA  NM_021800 Homo sapiens J domain containing protein 1 (JDP1), mRNA  NM_020404 Homo sapiens tumor endothelial marker 1 precursor (TEM1), mRNA	iated 2
(IRTA2), mRNA  NM_000767 Homo sapiens cytochrome P450, subfamily IIB (phenobarbital-inducible polypeptide 6 (CYP2B6), mRNA  NM_020165 Homo sapiens postreplication repair protein hRAD18p (RAD18), mRNA  NM_021800 Homo sapiens B-factor, properdin (BF), mRNA  NM_021800 Homo sapiens J domain containing protein 1 (JDP1), mRNA  NM_020404 Homo sapiens tumor endothelial marker 1 precursor (TEM1), mRNA	rated 2
NM_000767 Homo sapiens cytochrome P450, subfamily IIB (phenobarbital-inducible polypeptide 6 (CYP2B6), mRNA  NM_020165 Homo sapiens postreplication repair protein hRAD18p (RAD18), mRNA  NM_001710 Homo sapiens B-factor, properdin (BF), mRNA  NM_021800 Homo sapiens J domain containing protein 1 (JDP1), mRNA  NM_020404 Homo sapiens tumor endothelial marker 1 precursor (TEM1), mRNA	
polypeptide 6 (CYP2B6), mRNA  NM_020165 Homo sapiens postreplication repair protein hRAD18p (RAD18), mRNA  NM_001710 Homo sapiens B-factor, properdin (BF), mRNA  NM_021800 Homo sapiens J domain containing protein 1 (JDP1), mRNA  NM_020404 Homo sapiens tumor endothelial marker 1 precursor (TEM1), mRNA	`
NM_020165 Homo sapiens postreplication repair protein hRAD18p (RAD18), mRNA  NM_001710 Homo sapiens B-factor, properdin (BF), mRNA  NM_021800 Homo sapiens J domain containing protein 1 (JDP1), mRNA  NM_020404 Homo sapiens tumor endothelial marker 1 precursor (TEM1), mRNA	e),
NM 001710 Homo sapiens B-factor, properdin (BF), mRNA NM 021800 Homo sapiens J domain containing protein 1 (JDP1), mRNA NM 020404 Homo sapiens tumor endothelial marker 1 precursor (TEM1), mRNA	A
NM 021800 Homo sapiens J domain containing protein 1 (JDP1), mRNA NM 020404 Homo sapiens tumor endothelial marker 1 precursor (TEM1), mRNA	4
NM_020404 Homo sapiens tumor endothelial marker 1 precursor (TEM1), mRNA	
1 NIM 1006677 I Homo consona colute comion family 22 (anamia anim transporter)	
NM_006672 Homo sapiens solute carrier family 22 (organic anion transporter), members (SLC22A7), mRNA	Der /
NM_006398 Homo sapiens diubiquitin (UBD), mRNA	
	DNIA
1 07	nkna
	72274
NM_001632 Homo sapiens alkaline phosphatase, placental (Regan isozyme) (ALPP), NM_030773 Homo sapiens beta tubulin 1, class VI (TUBB1), mRNA	, MKNA
	T A
7,	
1 3	IA.
	743.60
NM_007023 Homo sapiens cAMP-regulated guanine nucleotide exchange factor II (C GEFII), mRNA	CAMP-
<u> </u>	. 11
NM_006105 Homo sapiens Rap1 guanine-nucleotide-exchange factor directly activat (EPAC), mRNA	ea by cA
NM_005637 Homo sapiens synovial sarcoma translocation, chromosome 18 (SS18),	mRNA
NM 001213 Homo sapiens chromosome 1 open reading frame 1 (Clorf1), mRNA	
NM_002354 Homo sapiens tumor-associated calcium signal transducer 1 (TACSTD1	).
mRNA	.,,
NM_003492 Homo sapiens chromosome X open reading frame 12 (CXorf12), mRNA	4
NM_003797 Homo sapiens embryonic ectoderm development (EED), mRNA	
NM_032863 Homo sapiens hypothetical protein FLJ14927 (FLJ14927), mRNA	
NM_032813 Homo sapiens hypothetical protein FLJ14624 (FLJ14624), mRNA	
NM_032578 Homo sapiens myopalladin (FLJ14437), mRNA	
NM_032385 Homo sapiens chromosome 5 open reading frame 4 (C5orf4), mRNA	
NM_032239 Homo sapiens hypothetical protein FLJ23511 (FLJ23511), mRNA	
NM 032012 Homo sapiens chromosome 9 open reading frame 5 (C9orf5), mRNA	
NM_031922 Homo sapiens RALBP1 protein (LOC83859), mRNA	
NM_031890 Homo sapiens cat eye syndrome chromosome region, candidate 6 (CEC)	R6).
mRNA	,
NM_031456 Homo sapiens chromosome 17 open reading frame 1A (C17orf1A), mR	NA
NM_030944 Homo sapiens chromosome 15 open reading frame 5 (C15orf5), mRNA	
NM_030806 Homo sapiens chromosome 1 open reading frame 21 (Clorf21), mRNA	
NM_030790 Homo sapiens hypothetical protein CDA08 (CDA08), mRNA	
NM 018312 Homo sapiens chromosome 11 open reading frame 23 (C11orf23), mRN	JA
NM_024298 Homo sapiens malignant cell expression-enhanced gene/tumor progressi	
enhanc (LENG4), mRNA	•
NM_022458 Homo sapiens chromosome 7 open reading frame 2 (C7orf2), mRNA	
NM_022338 Homo sapiens chromosome 11 open reading frame 24 (C11orf24), mRN	ĪΑ

ND 4 022162	The second of th
NM_022163	Homo sapiens chromosome 15 open reading frame 4 (C15orf4), mRNA
NM_022107	Homo sapiens chromosome 6 open reading frame 9 (C6orf9), mRNA
NM_006781	Homo sapiens chromosome 6 open reading frame 10 (C6orf10), mRNA
NM_019895	Homo sapiens chromosome 3 open reading frame 4 (C3orf4), mRNA
NM_012265	Homo sapiens chromosome 22 open reading frame 3 (C22orf3), mRNA
NM_021254	Homo sapiens chromosome 21 open reading frame 59 (C21orf59), mRNA
NM_020645	Homo sapiens chromosome 11 open reading frame 14 (C11orf14), mRNA
NM_012112	Homo sapiens chromosome 20 open reading frame 1 (C20orf1), mRNA
NM_018555	Homo sapiens zinc finger protein 331; zinc finger protein 463 (ZNF361), mRNA
NM_019106	Homo sapiens septin 3 (SEPT3), mRNA
NM_020375	Homo sapiens chromosome 12 open reading frame 5 (C12orf5), mRNA
NM_020374	Homo sapiens chromosome 12 open reading frame 4 (C12orf4), mRNA
NM_020373	Homo sapiens chromosome 12 open reading frame 3 (C12orf3), mRNA
NM_020367	Homo sapiens chromosome 12 open reading frame 6 (C12orf6), mRNA
NM_020130	Homo sapiens chromosome 8 open reading frame 4 (C8orf4), mRNA
NM_019596	Homo sapiens chromosome 21 open reading frame 62 (C21orf62), mRNA
NM_019063	Homo sapiens chromosome 2 open reading frame 2 (C2orf2), mRNA
NM_018956	Homo sapiens chromosome 9 open reading frame 9 (C9orf9), mRNA
NM_017586	Homo sapiens chromosome 9 open reading frame 7 (C9orf7), mRNA
NM_018691	Homo sapiens chromosome 5 open reading frame 3 (C5orf3), mRNA
NM_006134	Homo sapiens chromosome 21 open reading frame 4 (C21orf4), mRNA
NM_016940	Homo sapiens chromosome 21 open reading frame 6 (C21orf6), mRNA
NM_017438	Homo sapiens chromosome 21 open reading frame 18 (C21orf18), mRNA
NM_013265	Homo sapiens chromosome 11 open reading frame2 (C11orf2), mRNA
NM_016190	Homo sapiens chromosome 1 open reading frame 10 (C1orf10), mRNA
NM_015927	Homo sapiens transforming growth factor beta 1 induced transcript 1 (TGFB1I1), mRNA
NM 016564	Homo sapiens BM88 antigen (BM88), mRNA
NM 016348	Homo sapiens chromosome 5 open reading frame 4 (C5orf4), mRNA
NM 014009	Homo sapiens immune dysregulation, polyendocrinopathy, enteropathy, X-
	linked (IPEX), mRNA
NM_015524	Homo sapiens chromosome 6 open reading frame 5 (C6orf5), mRNA
NM_006345	Homo sapiens chromosome 4 open reading frame 1 (C4orf1), mRNA
NM_015373	Homo sapiens chromosome 22 open reading frame 2 (C22orf2), mRNA
NM_014205	Homo sapiens chromosome 11 open reading frame 5 (C11orf5), mRNA
NM_012264	Homo sapiens chromosome 22 open reading frame 5 (C22orf5), mRNA
NM_012111	Homo sapiens chromosome 14 open reading frame 3 (C14orf3), mRNA
NM_007211	Homo sapiens chromosome 12 open reading frame 2 (C12orf2), mRNA
NM_007176	Homo sapiens chromosome 14 open reading frame 1 (C14orf1), mRNA
NM_006706	Homo sapiens TATA box binding protein (TBP)-associated factor, RNA
) D ( 00 00 00	polymerase II, S, 150kD (TAF2S), mRNA
NM_006382	Homo sapiens chromosome 17 open reading frame 1A (C17orf1A), mRNA
NM_005967	Homo sapiens NGFI-A binding protein 2 (EGR1 binding protein 2) (NAB2), mRNA
NM_005966	Homo sapiens NGFI-A binding protein 1 (EGR1 binding protein 1) (NAB1), mRNA
NM_005663	Homo sapiens Wolf-Hirschhorn syndrome candidate 2 (WHSC2), mRNA
NM 005491	Homo sapiens chromosome X open reading frame 6 (CXorf6), mRNA
NM 005128	Homo sapiens chromosome 21 open reading frame 5 (C21orf5), mRNA
NM_004928	Homo sapiens chromosome 21 open reading frame 2 (C21orf2), mRNA
NM_004894	Homo sapiens chromosome 14 open reading frame 2 (C14orf2), mRNA
NM 004872	Homo sapiens chromosome 1 open reading frame 8 (C1orf8), mRNA

DD ( 004500	TT 1: 0 1 (GY, C)
NM_004709	Homo sapiens chromosome X open reading frame 1 (CXorf1), mRNA
NM_004337	Homo sapiens chromosome 8 open reading frame 1 (C8orf1), mRNA
NM_004913	Homo sapiens chromosome 16 open reading frame 7 (C16orf7), mRNA
NM_000956	Homo sapiens prostaglandin E receptor 2 (subtype EP2), 53kD (PTGER2), mRNA
NM 001586	
NM 001585	Homo sapiens chromosome X open reading frame 2 (CXorf2), mRNA  Homo sapiens chromosome 22 open reading frame 1 (C22orf1), mRNA
NM 001214	Homo sapiens chromosome 16 open reading frame 3 (C16orf3), mRNA
NM 001584	Homo sapiens chromosome 11 open reading frame 8 (C11orf8), mRNA  Homo sapiens chromosome 11 open reading frame 8 (C11orf8), mRNA
NM 003475	Homo sapiens chromosome 11 open reading frame 13 (C11orf13), mRNA  Homo sapiens chromosome 11 open reading frame 13 (C11orf13), mRNA
NM 032496	
NM 007234	Homo sapiens rho-gtpase activating protein ARHGAP9 (ARHGAP9), mRNA Homo sapiens dynactin 3 (p22) (DCTN3), transcript variant 1, mRNA
NM 024348	Homo sapiens dynactin 3 (p22) (DCTN3), transcript variant 1, mRNA  Homo sapiens dynactin 3 (p22) (DCTN3), transcript variant 2, mRNA
NM_021246	<u> </u>
NWI_021240	Homo sapiens megakaryocyte-enhanced gene transcript 1 protein (MEGT1), mRNA
NM_013291	Homo sapiens cleavage and polyadenylation specific factor 1, 160kD subunit (CPSF1), mRNA
NM_014500	Homo sapiens HIV TAT specific factor 1 (HTATSF1), mRNA
NM_005567	Homo sapiens lectin, galactoside-binding, soluble, 3 binding protein (LGALS3BP), mRNA
NM 005711	Homo sapiens EGF-like repeats and discoidin I-like domains 3 (EDIL3), mRNA
NM 016593	Homo sapiens oxysterol 7alpha-hydroxylase (CYP39A1), mRNA
NM 021048	Homo sapiens melanoma antigen, family A, 10 (MAGEA10), mRNA
NM 021049	Homo sapiens melanoma antigen, family A, 5 (MAGEA5), mRNA
NM 019602	Homo sapiens butyrophilin-like 2 (MHC class II associated) (BTNL2), mRNA
NM 018002	Homo sapiens oxidation resistance 1 (OXR1), mRNA
NM 013392	Homo sapiens nuclear receptor binding protein (NRBP), mRNA
NM_012396	Homo sapiens pleckstrin homology-like domain, family A, member 3
	(PHLDA3), mRNA
NM_006492	Homo sapiens aristaless-like homeobox 3 (ALX3), mRNA
NM_005365	Homo sapiens melanoma antigen, family A, 9 (MAGEA9), mRNA
NM_005364	Homo sapiens melanoma antigen, family A, 8 (MAGEA8), mRNA
NM_005366	Homo sapiens melanoma antigen, family A, 11 (MAGEA11), mRNA
NM_024490	Homo sapiens ATPase, Class V, type 10C (ATP10C), mRNA
NM_020354	Homo sapiens lysosomal apyrase-like protein 1 (LALP1), mRNA
NM_018655	Homo sapiens lens epithelial protein (LENEP), mRNA
NM_016448	Homo sapiens RA-regulated nuclear matrix-associated protein (RAMP), mRNA
NM_014763	Homo sapiens mitochondrial ribosomal protein L19 (MRPL19), mRNA
NM_006099	Homo sapiens protein inhibitor of activated STAT3 (PIAS3), mRNA
NM_004221	Homo sapiens natural killer cell transcript 4 (NK4), mRNA
NM_002949	Homo sapiens mitochondrial ribosomal protein L12 (MRPL12), mRNA
NM_016239	Homo sapiens myosin XVA (MYO15A), mRNA
NM_005094	Homo sapiens solute carrier family 27 (fatty acid transporter), member 4
	(SLC27A4), mRNA
NM_015077	Homo sapiens sterile alpha and HEAT/Armadillo motif protein, ortholog of Drosophila (SARM), mRNA
NM 013239	Homo sapiens protein phosphatase 2A 48 kDa regulatory subunit (PR48), mRNA
NM 022363	Homo sapiens LIM homeobox protein 5 (LHX5), mRNA
NM_031966	Homo sapiens cyclin B1 (CCNB1), mRNA
NM_015559	Homo sapiens SET binding protein 1 (SETBP1), mRNA
NM_007178	Homo sapiens unr-interacting protein (UNRIP), mRNA
NM_005367	Homo sapiens melanoma antigen, family A, 12 (MAGEA12), mRNA
1111 000001	1 1101110 Sapicits illetationia antigen, family A, 12 (MACEA12), Illetina

ND4 021275	Homo sapiens testis expressed sequence 12 (TEX12), mRNA
NM_031275	II a resistant mattered begin gamma subfamily C 2 (DCDUGC2) transcript
NM_032403	Homo sapiens protocadherin gamma subfamily C, 3 (PCDHGC3), transcript variant 3, mRNA
NM_032402	Homo sapiens protocadherin gamma subfamily C, 3 (PCDHGC3), transcript variant 2, mRNA
NM_002588	Homo sapiens protocadherin gamma subfamily C, 3 (PCDHGC3), transcript variant 1, mRNA
NM 014583	Homo sapiens LIM and cysteine-rich domains 1 (LMCD1), mRNA
	Homo sapiens Down syndrome cell adhesion molecule (DSCAM), mRNA
NM_001389	Homo sapiens ferritin, heavy polypeptide-like 17 (FTHL17), mRNA
NM_031894	Homo sapiens protocadherin gamma subfamily B, 4 (PCDHGB4), transcript
NM_032098	variant 2, mRNA
NM_003736	Homo sapiens protocadherin gamma subfamily B, 4 (PCDHGB4), transcript variant 1, mRNA
NM_032938	Homo sapiens G protein pathway suppressor 2 (GPS2), transcript variant 3, mRNA
NM_004489	Homo sapiens G protein pathway suppressor 2 (GPS2), transcript variant 2, mRNA
NM_032442	Homo sapiens G protein pathway suppressor 2 (GPS2), transcript variant 1, mRNA
NM 001887	Homo sapiens crystallin, beta B1 (CRYBB1), mRNA
NM 005208	Homo sapiens crystallin, beta A1 (CRYBA1), mRNA
NM 001889	Homo sapiens crystallin, zeta (quinone reductase) (CRYZ), mRNA
NM_022132	Homo sapiens methylcrotonoyl-Coenzyme A carboxylase 2 (beta) (MCCC2), mRNA
NM 001288	Homo sapiens chloride intracellular channel 1 (CLIC1), mRNA
NM 021624	Homo sapiens histamine H4 receptor (HRH4), mRNA
NM 032527	Homo sapiens hypothetical protein FLJ14972 (KIAA1847), mRNA
NM 005560	Homo sapiens laminin, alpha 5 (LAMA5), mRNA
NM 032931	Homo sapiens hypothetical protein MGC13219 (MGC13219), mRNA
NM 032924	Homo sapiens hypothetical protein MGC16040 (MGC16040), mRNA
NM 032920	Homo sapiens hypothetical protein MGC15873 (MGC15873), mRNA
NM 032913	Homo sapiens hypothetical protein MGC14458 (MGC14458), mRNA
NM_032893	Homo sapiens hypothetical protein MGC14336 (MGC14336), mRNA
NM 032889	Homo sapiens hypothetical protein MGC11308 (MGC11308), mRNA
NM 032815	Homo sapiens hypothetical protein FLJ14639 (FLJ14639), mRNA
NM 032798	Homo sapiens hypothetical protein FLJ14503 (FLJ14503), mRNA
NM 032793	Homo sapiens hypothetical protein FLJ14490 (FLJ14490), mRNA
NM 032791	Homo sapiens hypothetical protein FLJ14477 (FLJ14477), mRNA
NM 032789	Homo sapiens hypothetical protein FLJ14464 (FLJ14464), mRNA
NM 032769	Homo sapiens hypothetical protein MGC16212 (MGC16212), mRNA
NM 032760	Homo sapiens hypothetical protein MGC14966 (MGC14966), mRNA
NM 032696	Homo sapiens hypothetical protein MGC12262 (MGC12262), mRNA
NM 032665	Homo sapiens hypothetical protein MGC4640 (MGC4640), mRNA
NM 032662	Homo sapiens hypothetical protein MGC10600 (MGC10600), mRNA
NM 032655	Homo sapiens hypothetical protein MGC10997 (MGC10997), mRNA
NM 032625	Homo sapiens hypothetical brain protein my040 (MY040), mRNA
NM 032621	Homo sapiens X-linked protein (DJ79P11.1), mRNA
NM 032525	Homo sapiens tubulin beta-5 (TUBB5), mRNA
NM_005485	Homo sapiens ADP-ribosyltransferase (NAD+; poly (ADP-ribose) polymerase)-
NM_005484	like 3 (ADPRTL3), mRNA  Homo sapiens ADP-ribosyltransferase (NAD+; poly(ADP-ribose) polymerase)-

F	like 2 (ADPRTL2), mRNA
NM 005447	
14141_005447	Homo sapiens peptidylglycine alpha-amidating monooxygenase COOH-terminal interactor (PAMCI), mRNA
NM 000137	Homo sapiens fumarylacetoacetate hydrolase (fumarylacetoacetase) (FAH),
1.2.2_000107	mRNA
NM 001888	Homo sapiens crystallin, mu (CRYM), mRNA
NM 032608	Homo sapiens hypothetical protein bk125H2.1 (BK125H2.1), mRNA
NM 032607	Homo sapiens CREB/ATF family transcription factor (CREB-H), mRNA
NM 032602	Homo sapiens connexin 62 (CX62), mRNA
NM 032598	Homo sapiens testes development-related NYD-SP20 (NYD-SP20), mRNA
NM 032592	Homo sapiens 1-aminocyclopropane-1-carboxylate synthase (PHACS), mRNA
NM 032581	Homo sapiens down-regulated by Ctnnb1, a (DRCTNNB1A), mRNA
NM 032579	Homo sapiens colon and small intestine-specific cysteine-rich protein precursor
1111_0025,5	similar to FIZZ2/resistin-like protein (HXCP2), mRNA
NM 032570	Homo sapiens NPC-related protein NAG73 (NAG73), mRNA
NM 032565	Homo sapiens emopamil binding related protein, delta8-delta7 sterol isomerase
1111_032303	related protein (EBRP), mRNA
NM 032561	Homo sapiens EVG1 protein (EVG1), mRNA
NM 032555	Homo sapiens P143 protein (P143), mRNA
NM 032549	Homo sapiens inner mitochondrial membrane peptidase 2 like (IMMP2L),
1111_032349	mRNA
NM 032548	Homo sapiens BPOZ protein (BPOZ), mRNA
NM 015080	Homo sapiens neurexin 2 (NRXN2), mRNA
NM 005676	Homo sapiens RNA binding motif protein 10 (RBM10), mRNA
NM 032526	Homo sapiens cytosolic nucleotidase I (CN-I), mRNA
NM_032483	
NM 032094	Homo sapiens HTPAP protein (HTPAP), mRNA
14141_032094	Homo sapiens protocadherin gamma subfamily A, 12 (PCDHGA12), transcript variant 2, mRNA
NM_003735	Homo sapiens protocadherin gamma subfamily A, 12 (PCDHGA12), transcript variant 1, mRNA
NM 031887	Homo sapiens pro-melanin-concentrating hormone-like 1 (PMCHL1), mRNA
NM 032461	Homo sapiens SPANX family, member B1 (SPANXB1), mRNA
NM 006986	Homo sapiens melanoma antigen, family D, 1 (MAGED1), mRNA
NM 005462	Homo sapiens melanoma antigen, family C, 1 (MAGEC1), mRNA
NM_002375	Homo sapiens microtubule-associated protein 4 (MAP4), transcript variant 1,
	mRNA
NM_030983	Homo sapiens microtubule-associated protein 4 (MAP4), transcript variant 4, mRNA
NM 030885	
NM_030885	Homo sapiens microtubule-associated protein 4 (MAP4), transcript variant 3, mRNA
NM_030884	Homo sapiens microtubule-associated protein 4 (MAP4), transcript variant 2,
	mRNA
NM_002374	Homo sapiens microtubule-associated protein 2 (MAP2), transcript variant 1, mRNA
NM_031847	
	Homo sapiens microtubule-associated protein 2 (MAP2), transcript variant 4, mRNA
NM_031846	Homo sapiens microtubule-associated protein 2 (MAP2), transcript variant 3, mRNA
NM_031845	Homo sapiens microtubule-associated protein 2 (MAP2), transcript variant 2, mRNA
NM 032446	Homo sapiens MEGF10 protein (MEGF10), mRNA
NM_032417	Homo sapiens SPANX family, member D (SPANXD), mRNA

NM 013453	Homo sapiens sperm protein associated with the nucleus, X chromosome, family
_	member A1 (SPANXA1), mRNA
NM_020690	Homo sapiens KIAA1085 protein (KIAA1085), mRNA
NM_012121	Homo sapiens Cdc42 effector protein 4; binder of Rho GTPases 4 (CEP4),
	mRNA
NM_001019	Homo sapiens ribosomal protein S15a (RPS15A), mRNA
NM_022551	Homo sapiens ribosomal protein S18 (RPS18), mRNA
NM_005909	Homo sapiens microtubule-associated protein 1B (MAP1B), transcript variant 1, mRNA
NM_032010	Homo sapiens microtubule-associated protein 1B (MAP1B), transcript variant 2, mRNA
NM_002373	Homo sapiens microtubule-associated protein 1A (MAP1A), mRNA
NM_031366	Homo sapiens collagen, type IV, alpha 3 (Goodpasture antigen) (COL4A3), transcript variant 6, mRNA
NM_031365	Homo sapiens collagen, type IV, alpha 3 (Goodpasture antigen) (COL4A3), transcript variant 5, mRNA
NM_031364	Homo sapiens collagen, type IV, alpha 3 (Goodpasture antigen) (COL4A3), transcript variant 4, mRNA
NM_031363	Homo sapiens collagen, type IV, alpha 3 (Goodpasture antigen) (COL4A3), transcript variant 3, mRNA
NM_031362	Homo sapiens collagen, type IV, alpha 3 (Goodpasture antigen) (COL4A3), transcript variant 2, mRNA
NM_000091	Homo sapiens collagen, type IV, alpha 3 (Goodpasture antigen) (COL4A3), transcript variant 1, mRNA
NM_002140	Homo sapiens heterogeneous nuclear ribonucleoprotein K (HNRPK), transcript variant 1, mRNA
NM_031263	Homo sapiens heterogeneous nuclear ribonucleoprotein K (HNRPK), transcript variant 3, mRNA
NM_031262	Homo sapiens heterogeneous nuclear ribonucleoprotein K (HNRPK), transcript variant 2, mRNA
NM_032414	Homo sapiens prokineticin 1 precursor (PROK1), mRNA
NM_003214	Homo sapiens TEA domain family member 3 (TEAD3), mRNA
NM_015613	Homo sapiens DKFZP434K091 protein (PAL), mRNA
NM 030643	Homo sapiens apolipoprotein L, 4 (APOL4), mRNA
NM_022064	Homo sapiens hypothetical protein FLJ12565 (FLJ12565), mRNA
NM 017971	Homo sapiens mitochondrial ribosomal protein L20 (MRPL20), mRNA
NM_016504	Homo sapiens mitochondrial ribosomal protein L27 (MRPL27), mRNA
NM_014050	Homo sapiens mitochondrial ribosomal protein L42 (MRPL42), mRNA
NM 000014	Homo sapiens alpha-2-macroglobulin (A2M), mRNA
NM 004891	Homo sapiens mitochondrial ribosomal protein L33 (MRPL33), mRNA
NM 004864	Homo sapiens prostate differentiation factor (PLAB), mRNA
NM_000454	Homo sapiens superoxide dismutase 1, soluble (amyotrophic lateral sclerosis 1 (adult)) (SOD1), mRNA
NM 032391	Homo sapiens small nuclear protein PRAC (PRAC), mRNA
NM 032382	Homo sapiens hypothetical protein FLJ22315 (FLJ22315), mRNA
NM 032365	Homo sapiens hypothetical protein MGC5254 (MGC5254), mRNA
NM 032363	Homo sapiens HEIL2 protein (HEIL2), mRNA
NM_032335	Homo sapiens hypothetical protein MGC14797 (MGC14797), mRNA
NM_032276	Homo sapiens hypothetical protein DKFZp547E052 (DKFZp547E052), mRNA
NM_032272	Homo sapiens hypothetical protein DKFZp586G1123 (DKFZp586G1123), mRNA
NM_032260	Homo sapiens hypothetical protein DKFZp434P144 (DKFZp434P144), mRNA

NM_032237	Homo sapiens hypothetical protein FLJ23356 (FLJ23356), mRNA
NM_032220	Homo sapiens hypothetical protein FLJ22283 (FLJ22283), mRNA
NM_032219	Homo sapiens hypothetical protein FLJ22269 (FLJ22269), mRNA
NM_032204	Homo sapiens hypothetical protein FLJ21588 (FLJ21588), mRNA
NM_032203	Homo sapiens hypothetical protein FLJ21423 (FLJ21423), mRNA
NM_032202	Homo sapiens hypothetical protein FLJ21404 (FLJ21404), mRNA
NM 032173	Homo sapiens hypothetical protein FLJ12747 (FLJ12747), mRNA
NM 032157	Homo sapiens hypothetical protein FLJ11531 (FLJ11531), mRNA
NM_032150	Homo sapiens hypothetical protein DKFZp434P1735 (DKFZP434P1735), mRNA
NM_021005	Homo sapiens nuclear receptor subfamily 2, group F, member 2 (NR2F2), mRNA
NM_020159	Homo sapiens hypothetical protein DKFZp762K2015 (DKFZp762K2015), mRNA
NM_015449	Homo sapiens DKFZP586G1722 protein (DKFZP586G1722), mRNA
NM_015424	Homo sapiens DKFZP586N2124 protein (DKFZP586N2124), mRNA
NM_015235	Homo sapiens likely ortholog of mouse variant polyadenylation protein CSTF-64; KIAA0689 protein (KIAA0689), mRNA
NM 015068	Homo sapiens paternally expressed 10 (PEG10), mRNA
NM 014599	Homo sapiens EH-domain containing 4 (EHD4), mRNA
NM_014411	Homo sapiens brain and nasopharyngeal carcinoma susceptibility protein (NSG-X), mRNA
NM 007148	Homo sapiens zinc finger protein 179 (ZNF179), mRNA
NM_007266	Homo sapiens XPA binding protein 1; putative ATP(GTP)-binding protein (NTPBP), mRNA
NM 006313	Homo sapiens ubiquitin specific protease 15 (USP15), mRNA
NM 005726	Homo sapiens Ts translation elongation factor, mitochondrial (TSFM), mRNA
NM 005277	Homo sapiens glycoprotein M6A (GPM6A), mRNA
NM 005437	Homo sapiens nuclear receptor coactivator 4 (NCOA4), mRNA
NM 001439	Homo sapiens exostoses (multiple)-like 2 (EXTL2), mRNA
NM 001287	Homo sapiens chloride channel 7 (CLCN7), mRNA
NM_021194	Homo sapiens solute carrier family 30 (zinc transporter), member 1 (SLC30A1), mRNA
NM_013986	Homo sapiens Ewing sarcoma breakpoint region 1 (EWSR1), transcript variant EWS-b, mRNA
NM 001013	Homo sapiens ribosomal protein S9 (RPS9), mRNA
NM 005617	Homo sapiens ribosomal protein S14 (RPS14), mRNA
NM 006361	Homo sapiens homeo box B13 (HOXB13), mRNA
NM 000990	Homo sapiens ribosomal protein L27a (RPL27A), mRNA
NM 005821	Homo sapiens NBR2 (NBR2), mRNA
NM_003483	Homo sapiens high-mobility group (nonhistone chromosomal) protein isoform I-C (HMGIC), mRNA
NM_002129	Homo sapiens high-mobility group (nonhistone chromosomal) protein 2 (HMG2), mRNA
NM 005959	Homo sapiens melatonin receptor 1B (MTNR1B), mRNA
NM 005958	Homo sapiens melatonin receptor 1A (MTNR1A), mRNA
NM 004739	Homo sapiens metastasis-associated 1-like 1 (MTA1L1), mRNA
NM 021644	Homo sapiens heterogeneous nuclear ribonucleoprotein H3 (2H9) (HNRPH3),
	transcript variant 2H9A, mRNA
NM_012207	Homo sapiens heterogeneous nuclear ribonucleoprotein H3 (2H9) (HNRPH3), transcript variant 2H9, mRNA
NM 019597	Homo sapiens heterogeneous nuclear ribonucleoprotein H2 (H') (HNRPH2),
	(-, (-, -, -, -, -, -, -, -, -, -, -, -, -, -

mRNA Homo sapiens heterogeneous nuclear ribonucleoprotein M (HNRPM), transcript variant 2, mRNA Homo sapiens heterogeneous nuclear ribonucleoprotein M (HNRPM), transcript
variant 2, mRNA
Transcript and the following induction in the same of
variant 1, mRNA
Homo sapiens heterogeneous nuclear ribonucleoprotein F (HNRPF), mRNA
Homo sapiens pregnancy-associated interferon (HTIFN), mRNA
Homo sapiens mitochondrial ribosomal protein L1 (MRPL1), mRNA
Homo sapiens mitochondrial ribosomal protein L11 (MRPL11), mRNA
Homo sapiens heterogeneous nuclear ribonucleoprotein H1 (H) (HNRPH1),
mRNA
Homo sapiens jagged 2 (JAG2), mRNA
Homo sapiens heterogeneous nuclear ribonucleoprotein A0 (HNRPA0), mRNA
Homo sapiens heterogeneous nuclear ribonucleoprotein D-like (HNRPDL),
transcript variant 1, mRNA
Homo sapiens heterogeneous nuclear ribonucleoprotein D-like (HNRPDL),
transcript variant 2, mRNA
Homo sapiens alkaline phosphatase, placental-like 2 (ALPPL2), mRNA
Homo sapiens X-box binding protein 1 (XBP1), mRNA
Homo sapiens cell division cycle 2-like 5 (cholinesterase-related cell division
controller) (CDC2L5), transcript variant 2, mRNA
Homo sapiens cell division cycle 2-like 5 (cholinesterase-related cell division
controller) (CDC2L5), transcript variant 1, mRNA
Homo sapiens cytochrome P450, subfamily IID (debrisoquine, sparteine, etc., -
metabolizing), polypeptide 6 (CYP2D6), mRNA
Homo sapiens membrane component, chromosome 17, surface marker 2 (ovarian
carcinoma antigen CA125) (M17S2), transcript variant 3, mRNA
Homo sapiens membrane component, chromosome 17, surface marker 2 (ovarian
carcinoma antigen CA125) (M17S2), transcript variant 2, mRNA
Homo sapiens membrane component, chromosome 17, surface marker 2 (ovarian
carcinoma antigen CA125) (M17S2), transcript variant 1, mRNA
Homo sapiens hypothetical protein DKFZp547N043 (DKFZP547N043), mRNA
Homo sapiens testes-specific heterogenous nuclear ribonucleoprotein G-T
(HNRNPG-T), mRNA
Homo sapiens heterogeneous nuclear ribonucleoprotein A2/B1 (HNRPA2B1),
transcript variant A2, mRNA
Homo sapiens heterogeneous nuclear ribonucleoprotein A2/B1 (HNRPA2B1),
transcript variant B1, mRNA
Homo sapiens heterogeneous nuclear ribonucleoprotein A1 (HNRPA1),
transcript variant 2, mRNA
Homo sapiens angiotensin receptor 1 (AGTR1), transcript variant 2, mRNA
Homo sapiens angiotensin receptor 1 (AGTR1), transcript variant 5, mRNA
Homo sapiens angiotensin receptor 1 (AGTR1), transcript variant 4, mRNA
Homo sapiens angiotensin receptor 1 (AGTR1), transcript variant 3, mRNA
Homo sapiens angiotensin receptor 1 (AGTR1), transcript variant 1, mRNA
Homo sapiens chemokine (C-C motif) receptor-like 2 (CCRL2), mRNA
Homo sapiens chemokine (C-C motif) receptor 9 (CCR9), transcript variant B, mRNA
Homo sapiens chemokine (C-C motif) receptor 9 (CCR9), transcript variant A,
mRNA
Homo sapiens chemokine (C-C motif) receptor 6 (CCR6), transcript variant 2,
mRNA

NM_004367	Homo sapiens chemokine (C-C motif) receptor 6 (CCR6), transcript variant 1, mRNA
NM 031371	Homo sapiens RBP1-like protein (BCAA), transcript variant 2, mRNA
NM 016374	Homo sapiens RBP1-like protein (BCAA), transcript variant 2, mRNA
NM 004281	Homo sapiens BCL2-associated athanogene 3 (BAG3), mRNA
NM 032048	Homo sapiens extracellular glycoprotein EMILIN-2 precursor (EMILIN-2),
	mRNA
NM_032046	Homo sapiens mosaic serine protease (MSP), mRNA
NM_032045	Homo sapiens kringle-containing transmembrane protein; kringle-coding gene marking the eye and the nose (KREMEN), mRNA
NM 032044	Homo sapiens regenerating gene type IV (REG-IV), mRNA
NM 032041	Homo sapiens neurocalcin delta (NCALD), mRNA
NM_032039	Homo sapiens hypothetical protein DKFZp761D0211 (DKFZP761D0211), mRNA
NM_032038	Homo sapiens spinster-like protein (LOC83985), mRNA
NM_032020	Homo sapiens hypothetical protein MGC1314 similar to fucosidase, alpha-L-1,
	Tissue (MGC1314), mRNA
NM_032016	Homo sapiens hypothetical protein MGC3251 (MGC3251), mRNA
NM_000323	Homo sapiens ret proto-oncogene (multiple endocrine neoplasia and medullary
) D 6 6555	thyroid carcinoma 1, Hirschsprung disease) (RET), transcript variant 1, mRNA
NM_020975	Homo sapiens ret proto-oncogene (multiple endocrine neoplasia and medullary
) The 600 co	thyroid carcinoma 1, Hirschsprung disease) (RET), transcript variant 2 mRNA
NM_020630	Homo sapiens ret proto-oncogene (multiple endocrine neoplasia and medullary
) N 6 000 600	inyroid carcinoma I, Hirschsprung disease) (RET), transcript variant 4 mRNA
NM_020629	Homo sapiens ret proto-oncogene (multiple endocrine neoplasia and medullary)
ND 6 01 6015	thyroid carcinoma 1, Hirschsprung disease) (RET), transcript variant 3, mRNA
NM_016817	Homo sapiens 2'-5'-oligoadenylate synthetase 2 (69-71 kD) (OAS2), transcript variant 1, mRNA
NM 006187	Homo sapiens 2'-5'-oligoadenylate synthetase 3 (100 kD) (OAS3), mRNA
NM_002535	Homo sapiens 2'-5'-oligoadenylate synthetase 2 (69-71 kD) (OAS2), transcript
	variant 2, mRNA
NM_002342	Homo sapiens lymphotoxin beta receptor (TNFR superfamily, member 3) (LTBR), mRNA
NM_002136	Homo sapiens heterogeneous nuclear ribonucleoprotein A1 (HNRPA1),
	transcript variant 1, mRNA
NM_001885	Homo sapiens crystallin, alpha B (CRYAB), mRNA
NM_015139	Homo sapiens UDP-glucuronic acid/UDP-N-acetylgalactosamine dual
	transporter (UGTREL7), mRNA
NM_024333	Homo sapiens fibronectin type 3 and SPRY domain-containing protein (FSD1),
NM_017947	mRNA Homo conjone melalala Control (Control (Con
NM_017934	Homo sapiens molybdenum cofactor sulfurase (HMCS), mRNA
NM 016492	Homo sapiens pleckstrin homology domain interacting protein (PHIP), mRNA
NM_014185	Homo sapiens homolog of yeast MOG1 (MOG1), mRNA
NM 031965	Homo sapiens homolog of yeast MOG1 (MOG1), mRNA
NM 031952	Homo sapiens haspin (GSG2), mRNA
NM 031950	Homo sapiens NYD-SP16 protein (NYD-SP16), mRNA
NM_031949	Homo sapiens NVD TSDG (KSP37), mRNA
NM_031945	Homo sapiens NYD-TSPG protein (NYD-TSPG), mRNA
NM_031943	Homo sapiens (ED28 (ED28), mRNA
NM_031943	Homo sapiens (Mrs. 1974), mRNA
NM 031942	Homo sapiens AJE 751: 1: 040000 PN
11111_031341	Homo sapiens AIE-75 binding protein protein (MCC2), mRNA

	TI
NM_031938	Homo sapiens putative b,b-carotene-9',10'-dioxygenase (B-DIOX-II), mRNA
NM_031937	Homo sapiens EBP50-PDZ interactor of 64 kD (EPI64), mRNA
NM_031921	Homo sapiens AAA-ATPase TOB3 (TOB3), mRNA
NM_031915	Homo sapiens CLLL8 protein (CLLD8), mRNA
NM_031911	Homo sapiens complement-c1q tumor necrosis factor-related protein 7 (CTRP7), mRNA
NM_031910	Homo sapiens complement-clq tumor necrosis factor-related protein 6 (CTRP6), mRNA
NM_031909	Homo sapiens complement-c1q tumor necrosis factor-related protein 4 (CTRP4), mRNA
NM 031904	Homo sapiens hypothetical protein FKSG44 (FKSG44), mRNA
NM 031903	Homo sapiens mitochondrial ribosomal protein L32 (MRPL32), mRNA
NM 031900	Homo sapiens alanine-glyoxylate aminotransferase 2 (AGXT2), mRNA
NM 031897	Homo sapiens calcium channel, voltage-dependent, gamma subunit 6
_	(CACNG6), mRNA
NM_031896	Homo sapiens calcium channel, voltage-dependent, gamma subunit 7 (CACNG7), mRNA
NM 031939	Homo sapiens B29 protein (B29), mRNA
NM_031886	Homo sapiens potassium voltage-gated channel, shaker-related subfamily, member 7 (KCNA7), mRNA
NM 020992	Homo saniens PDZ and LIM domain 1 (elfin) (PDLIM1), mRNA
NM 031407	Homo sapiens upstream regulatory element binding protein 1 (UREB1), mRNA
NM 030582	Homo sapiens collagen, type XVIII, alpha 1 (COL18A1), mRNA
NM 020390	Homo sapiens eukaryotic translation initiation factor 5A2 (EIF5A2), mRNA
NM 018980	Homo sapiens taste receptor, type 2, member 5 (TAS2R5), mRNA
NM 018417	Homo sapiens soluble adenylyl cyclase (SAC), mRNA
NM 016945	Homo sapiens taste receptor, type 2, member 16 (TAS2R16), mRNA
NM_004775	Homo sapiens UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 6 (B4GALT6), mRNA
NM_003778	Homo sapiens UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 4 (B4GALT4), mRNA
NM_003779	Homo sapiens UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 3 (B4GALT3), mRNA
NM 001296	Homo sapiens chemokine binding protein 2 (CCBP2), mRNA
NM 001497	Homo sapiens UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide
_	1 (B4GALT1), mRNA
NM 014451	Homo sapiens PTH-responsive osteosarcoma B1 protein (B1), mRNA
NM_031265	Homo sapiens mucin and cadherin-like (MUCDHL), transcript variant 4, mRNA
NM 031264	Homo sapiens mucin and cadherin-like (MUCDHL), transcript variant 3, mRNA
NM 017717	Homo sapiens mucin and cadherin-like (MUCDHL), transcript variant 2, mRNA
NM 021924	Homo sapiens mucin and cadherin-like (MUCDHL), transcript variant 1, mRNA
NM 019855	Homo sapiens calcium binding protein 5 (CABP5), mRNA
NM_016367	Homo sapiens calcium binding protein 3 (CABP3), mRNA
NM_031204	Homo sapiens calcium binding protein 2 (CABP2), transcript variant 2, mRNA
NM_005201	Homo sapiens chemokine (C-C motif) receptor 8 (CCR8), mRNA
NM_000786	Homo sapiens cytochrome P450, 51 (lanosterol 14-alpha-demethylase) (CYP51), mRNA
NM_030908	Homo sapiens olfactory receptor, family 2, subfamily A, member 4 (OR2A4), mRNA
NM 001009	Homo sapiens ribosomal protein S5 (RPS5), mRNA
NM 001032	Homo sapiens ribosomal protein S29 (RPS29), mRNA
NM 001014	Homo sapiens ribosomal protein S10 (RPS10), mRNA
	1

NM 000991	Homo sapiens ribosomal protein L28 (RPL28), mRNA
NM 000782	Homo sapiens cytochrome P450, subfamily XXIV (vitamin D 24-hydroxylase)
11212_000702	(CYP24), mitochondrial protein encoded by nuclear gene, mRNA
NM 031226	Homo sapiens cytochrome P450, subfamily XIX (aromatization of androgens)
	(CYP19), transcript variant 2, mRNA
NM 000103	Homo sapiens cytochrome P450, subfamily XIX (aromatization of androgens)
	(CYP19), transcript variant 1, mRNA
NM 000498	Homo sapiens cytochrome P450, subfamily XIB (steroid 11-beta-hydroxylase),
_	polypeptide 2 (CYP11B2), mitochondrial protein encoded by nuclear gene,
	mRNA
NM 000102	Homo sapiens cytochrome P450, subfamily XVII (steroid 17-alpha-
_	hydroxylase), adrenal hyperplasia (CYP17), mRNA
NM_000497	Homo sapiens cytochrome P450, subfamily XIB (steroid 11-beta-hydroxylase),
_	polypeptide 1 (CYP11B1), mitochondrial protein encoded by nuclear gene,
	mRNA
NM_017460	Homo sapiens cytochrome P450, subfamily IIIA (niphedipine oxidase),
	polypeptide 4 (CYP3A4), mRNA
NM_018482	Homo sapiens development and differentiation enhancing factor 1 (DDEF1),
	mRNA
NM_016366	Homo sapiens calcium binding protein 2 (CABP2), transcript variant 1, mRNA
NM_007255	Homo sapiens xylosylprotein beta 1,4-galactosyltransferase, polypeptide 7
	(galactosyltransferase I) (B4GALT7), mRNA
NM_006668	Homo sapiens cytochrome P450, subfamily 46 (cholesterol 24-hydroxylase)
	(CYP46), mRNA
NM_000781	Homo sapiens cytochrome P450, subfamily XIA (cholesterol side chain
	cleavage) (CYP11A), nuclear gene encoding mitochondrial protein, mRNA
NM_000579	Homo sapiens chemokine (C-C motif) receptor 5 (CCR5), mRNA
NM_001295	Homo sapiens chemokine (C-C motif) receptor 1 (CCR1), mRNA
NM_031492	Homo sapiens hypothetical protein similar to RNA-binding protein lark (MGC10871), mRNA
NM_031488	Homo sapiens hypothetical protein DKFZp761I141 (DKFZP761I141), mRNA
NM_031469	Homo sapiens SH3 domain binding glutamic acid-rich protein like 2
	(SH3BGRL2), mRNA
NM_031468	Homo sapiens calneuron 1 (CALN1), mRNA
NM_031462	Homo sapiens hypothetical protein DKFZp761H2024 (DKFZP761H2024),
	mRNA
NM_031458	Homo sapiens B aggressive lymphoma gene (BAL), mRNA
NM_031445	Homo sapiens hypothetical protein MGC4268 (MGC4268), mRNA
NM_031440	Homo sapiens transmembrane protein 7 (TMEM7), mRNA
NM_031429	Homo sapiens retbindin (RTBDN), mRNA
NM_031427	Homo sapiens hypothetical protein MGC12435 (MGC12435), mRNA
NM_031426	Homo sapiens hypothetical protein FLJ12783 (FLJ12783), mRNA
NM_031422	Homo sapiens GalNAc-4-sulfotransferase 2 (GALNAC4ST-2), mRNA
NM_031415	Homo sapiens melanoma-derived leucine zipper, extra-nuclear factor (MLZE),
NM 031413	mRNA
U31413	Homo sapiens cat eye syndrome chromosome region, candidate 2 (CECR2), mRNA
NM_022719	Homo sapiens DiGeorge syndrome critical region gene DGSI; likely ortholog of
	mouse expressed sequence 2 embryonic lethal (DGSI), mRNA
NM_000669	Homo sapiens alcohol dehydrogenase 1C (class I), gamma polypeptide
_	(ADH1C), mRNA

	mRNA
NM_018833	Homo sapiens transporter 2, ATP-binding cassette, sub-family B (MDR/TAP) (TAP2), transcript variant 2, mRNA
ND 6 000544	Homo sapiens transporter 2, ATP-binding cassette, sub-family B (MDR/TAP)
NM_000544	Homo sapiens transporter 2, ATF-binding cassette, sub-taining B (1992)
	(TAP2), transcript variant 1, mRNA  Homo sapiens transporter 1, ATP-binding cassette, sub-family B (MDR/TAP)
NM_000593	
	(TAP1), mRNA
NM_004678	Homo sapiens variable charge, Y chromosome, 2 (VCY2), mRNA
NM_012392	Homo sapiens PEF protein with a long N-terminal hydrophobic domain (peflin) (PEF), mRNA
NM 031308	Homo sapiens epiplakin 1 (EPPK1), mRNA
NM 031299	Homo sapiens hypothetical protein MGC2577 (MGC2577), mRNA
NM 012480	Homo sapiens zinc finger protein 73 (Cos12) (ZNF73), mRNA
NM_030881	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 17 (72kD)
14147_030001	(DDX17), transcript variant 2, mRNA
NM_006386	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 17 (72kD)
14141_000390	(DDX17), transcript variant 1, mRNA
NM_003587	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 16 (DDX16),
	mRNA
NM 000478	Homo sapiens alkaline phosphatase, liver/bone/kidney (ALPL), mRNA
NM_004820	Homo sapiens cytochrome P450, subfamily VIIB (oxysterol 7 alpha-
<del>-</del>	hydroxylase), polypeptide 1 (CYP7B1), mRNA
NM 000780	Homo sapiens cytochrome P450, subfamily VIIA (cholesterol 7 alpha-
	monooxygenase), polypeptide 1 (CYP7A1), nuclear gene encoding
	mitochondrial protein, mRNA
NM 016166	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box binding protein 1
11111_010100	(DDXBP1), mRNA
NM 016373	Homo sapiens WW domain-containing oxidoreductase (WWOX), mRNA
NM 024164	Homo sapiens tryptase beta 2 (TPSB2), mRNA
NM 003294	Homo sapiens tryptase beta 1 (TPSB1), mRNA
NM 031310	Homo sapiens fenestrated-endothelial linked structure protein; PV-1 protein
14141_031310	(PV1), mRNA
NM 031302	Homo sapiens gycosyltransferase (LOC83468), mRNA
	Homo sapiens hypothetical protein MGC2383 (MGC2383), mRNA
NM_031300	Homo sapiens hypothetical protein DKFZp761H1710 (DKFZP761H1710),
NM_031297	
	mRNA Homo sapiens hypothetical protein MGC3133 (MGC3133), mRNA
NM_031287	Homo sapiens hypothetical protein MGC5155 (MGC5155), midva
NM_031286	Homo sapiens SH3BGRL3-like protein (SH3BGRL3), mRNA
NM_031285	Homo sapiens hypothetical protein PP1057 (PP1057), mRNA
NM_031279	Homo sapiens alanine-glyoxylate aminotransferase 2-like 1 (AGXT2L1), mRNA
NM_030970	Homo sapiens hypothetical protein MGC3771 (MGC3771), mRNA
NM_014357	Homo sapiens skin-specific protein (XP5), mRNA
NM_030590	Homo sapiens matrilin 4 (MATN4), transcript variant 2, mRNA
NM_031246	Homo sapiens pregnancy specific beta-1-glycoprotein 2 (PSG2), mRNA
NM 017422	Homo sapiens calmodulin-like skin protein (CLSP), mRNA
14141 O1/477	Homo sapiens methylenetetrahydrofolate dehydrogenase (NADP+ dependent),
NM_017422 NM_005956	methenyltetrahydrofolate cyclohydrolase, formyltetrahydrofolate synthetase
	methenyltetrahydrofolate cyclohydrolase, formyltetrahydrofolate synthetase (MTHFD1), mRNA
NM_005956	(MTHFD1), mRNA
NM_005956 NM_005906	(MTHFD1), mRNA  Homo sapiens male germ cell-associated kinase (MAK), mRNA
NM_005956	(MTHFD1), mRNA

	V) (TBXAS1), transcript variant TXS-II, mRNA
NM 001061	Homo sapiens thromboxane A synthase 1 (platelet, cytochrome P450, subfamily
NM_001001	V) (TBXAS1), transcript variant TXS-I, mRNA
ND ( 000772	Homo sapiens cytochrome P450, subfamily IIE (ethanol-inducible) (CYP2E),
NM_000773	mRNA
NM 030592	Homo sapiens matrilin 4 (MATN4), transcript variant 3, mRNA
NM_003833	Homo sapiens matrilin 4 (MATN4), transcript variant 1, mRNA
NM_005355	Homo sapiens kinesin-like 3 (KNSL3), transcript variant 2, mRNA
NM_030615	Homo sapiens kinesin-like 3 (KNSL3), transcript variant 1, mRNA
NM_004523	Homo sapiens kinesin-like 1 (KNSL1), mRNA
NM_005000	Homo sapiens NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 5
ND4 004541	(13kD, B13) (NDUFA5), nuclear gene encoding mitochondrial protein, mRNA
NM_004541	Homo sapiens NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 1
	(7.5kD, MWFE) (NDUFA1), nuclear gene encoding mitochondrial protein,
NB/ 000771	mRNA
NM_000771	Homo sapiens cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase),
ND4 000772	polypeptide 9 (CYP2C9), mRNA
NM_000772	Homo sapiens cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase),
NM 017778	polypeptide 18 (CYP2C18), mRNA  Homo sapiens Wolf-Hirschhorn syndrome candidate 1-like 1 (WHSC1L1),
NM_01///8	
NM 023034	transcript variant short, mRNA  Homo sapiens Wolf-Hirschhorn syndrome candidate 1-like 1 (WHSC1L1),
10101_023034	transcript variant long, mRNA
NM 000766	Homo sapiens cytochrome P450, subfamily IIA (phenobarbital-inducible),
NM_000766	polypeptide 13 (CYP2A13), mRNA
NM 006646	Homo sapiens WAS protein family, member 3 (WASF3), mRNA
NM 018560	Homo sapiens WW domain-containing oxidoreductase (WWOX), mRNA
NM_014110	Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 8 (PPP1R8),
11/11_014110	mRNA
NM_004109	Homo sapiens ferredoxin 1 (FDX1), nuclear gene encoding mitochondrial
11111_004105	protein, mRNA
NM 030671	Homo sapiens protein tyrosine phosphatase, receptor type, O (PTPRO),
11111_050071	transcript variant 5, mRNA
NM 030670	Homo sapiens protein tyrosine phosphatase, receptor type, O (PTPRO),
11212_00000	transcript variant 6, mRNA
NM 030669	Homo sapiens protein tyrosine phosphatase, receptor type, O (PTPRO),
	transcript variant 3, mRNA
NM 030668	Homo sapiens protein tyrosine phosphatase, receptor type, O (PTPRO),
	transcript variant 4, mRNA
NM 030667	Homo sapiens protein tyrosine phosphatase, receptor type, O (PTPRO),
	transcript variant 1, mRNA
NM 002848	Homo sapiens protein tyrosine phosphatase, receptor type, O (PTPRO),
_	transcript variant 2, mRNA
NM 021979	Homo sapiens heat shock 70kD protein 2 (HSPA2), mRNA
NM 024005	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 3 (DDX3),
_	transcript variant 1, mRNA
NM 001356	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 3 (DDX3),
_	transcript variant 2, mRNA
NM 020216	Homo sapiens arginyl aminopeptidase (aminopeptidase B) (RNPEP), mRNA
NM 006990	Homo sapiens WAS protein family, member 2 (WASF2), mRNA
NM 012467	Homo sapiens tryptase gamma 1 (TPSG1), mRNA
NM 007317	Homo sapiens kinesin-like 4 (KNSL4), mRNA

VM_004256	Homo sapiens organic cationic transporter-like 3 (ORCTL3), mRNA
VM_000774	Homo sapiens cytochrome P450, subfamily IIF, polypeptide 1 (CYP2F1),
	mRNA 15 15 15 HC (monhenytoin 4-hydroxylase).
VM_000769	Homo sapiens cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase),
	polypeptide 19 (CYP2C19), mRNA
NM 031220	Homo sapiens PYK2 N-terminal domain-interacting receptor 1 (NIR1), mRNA
NM 031212	Homo sapiens hypothetical protein NPD016 (NPD016), mRNA
NM 031211	Homo saniens I AT1-3TM protein (LAT1-3TM), mRNA
NM 031209	Homo saniens tRNA-guanine transglycosylase (TGT), mRNA
NM 031206	77 Is an atherical protein FI 112525 (F) (12525), IIIKINA
NM_006904	Homo sapiens protein kinase, DNA-activated, catalytic polypeptide (FREDE),
NM_030963	Homo sapiens hypothetical protein DKFZp434O1427 (DKFZP434O1427),
	mRNA (FSP13.2) mRNA
NM_030931	Homo sapiens epididymal secretory protein ESP13.2 (ESP13.2), mRNA  Homo sapiens epididymal secretory protein ESP13.2 (ESP13.2), mRNA
NM_030905	Homo sapiens olfactory receptor, family 2, subfamily J, member 2 (OR2J2), mRNA
NM_030903	Homo sapiens olfactory receptor, family 2, subfamily W, member 1 (OR2W1), mRNA
NM_012377	Homo sapiens olfactory receptor, family 7, subfamily C, member 2 (OR7C2), mRNA
ND 6 020001	Hamo copiens small GTP-hinding protein (RAB1B), mRNA
NM_030981	Homo sapiens smart GTT omening protein DKFZp434N1923 (DKFZP434N1923),
NM_030974	DNIA
NM 030973	Homo sapiens hypothetical protein TCBAP0758 (TCBAP0758), mRNA
NM_030968	Homo sapiens hypothetical protein 1932 the Homo sapiens G protein coupled receptor interacting protein, complement-clq tumor necrosis factor-related (ZSIG37), mRNA
NM_030945	Transport complement of a tumor necrosis factor-related protein; likely
14141_030543	ortholog of mouse CORS26 (collagenous repeat-containing sequence of 20-kba
37 f 020026	Homo sapiens hypothetical protein DKFZp434C135 (DKFZP434C135i), mRNA
NM_030936	Homo sapiens TSC-22-like (THG-1), mRNA
NM_030935	Homo sapiens integral membrane protein 3 (ITM3), mRNA
NM_030926	Homo sapiens integral memorane protein 5 (17145), many
NM_030893	Homo sapiens CD1E antigen, e polypeptide (CD1E), mRNA
NM_014067	Homo sapiens LRP16 protein (LRP16), mRNA
NM_030661	Homo sapiens homeo box A3 (HOXA3), mRNA
NM_030879	Homo sapiens Small evolutionarily conserved RNA, resembling C/D box small nucleolar (X102), mRNA
NM_012373	Homo sapiens olfactory receptor, family 3, subfamily A, member 3 (OR3A3), mRNA
NM 015072	Homo sapiens KIAA0998 protein (KIAA0998), mRNA
NM 030882	Homo sapiens apolipoprotein L, 2 (APOL2), mRNA
NM 002623	Homo sapiens prefoldin 4 (PFDN4), mRNA
NM 022167	Home conjens vylosyltransferase II (XT2), mRNA
NM 017506	
NM_017300	mPNA
NM 003372	Homo sapiens von Hippel-Lindau binding protein 1 (VBP1), mRNA
NM 016097	Homo sapiens HSPC039 protein (HSPC039), mRNA
NM_014646	Home seniens linin 2 (I PIN2) mRNA
NM_005880	Homo sapiens DnaJ (Hsp40) homolog, subfamily A, member 2 (DNAJA2), mRNA
NM 006755	TO A

	Homo sapiens DiGeorge syndrome critical region gene 2 (DGCR2), mRNA
NM_005137	Homo sapiens DiGeorge syndrome critical region gene 2 (DGCR2), mid 17
NM 000022	Homo sapiens adenosine deaminase (ADA), mRNA  Homo sapiens tec protein tyrosine kinase (TEC), mRNA
NM_003215	Homo sapiens the protein tyrosine khase (TEC), indext.  Homo sapiens phosphatidylinositol 4-kinase type II (PI4KII), mRNA
NM_018425	Homo sapiens BTB (POZ) domain containing 1 (BTBD1), mRNA
NM_025238	Homo sapiens G protein-coupled receptor 10 (GPR10), mRNA
NM_004248	Homo sapiens amyloid beta (A4) precursor-like protein 2 (APLP2), mRNA
NM_001642	Homo sapiens amyloid beta (A4) precursor-like protein 2 (A1 Li 2), ind (12)
NM_030821	Homo sapiens group XII secreted phospholipase A2 (PLA2G12), mRNA
NM_030820	Homo sapiens hypothetical protein DKFZp564B052 (DKFZp564B052), mRNA
NM_030816	Homo sapiens hypothetical protein DKFZp566D1346 (DKFZP566D1346), mRNA
NM 030807	Homo sapiens glucose transporter protein 10 (GLUT10), mRNA
NM_030798	Homo sapiens hypothetical protein DKFZp434D0421 (DKFZP434D0421), mRNA
NM_030797	Homo sapiens hypothetical protein DKFZp566A1524 (DKFZP566A1524), mRNA
NM 030788	Homo sapiens DC-specific transmembrane protein (LOC81501), mRNA
NM 030787	Homo sapiens factor H-related protein 5 (FHR5), mRNA
NM 030786	Homo sapiens intermediate filament protein syncoilin (SYNCOILIN), mRNA
NM 030785	Homo sapiens ortholog of mouse radial spokehead-like 1 (RSHL1), mRNA
NM 030784	Homo sapiens brain expressed G-protein-coupled receptor PSP24 beta
	(PSP24B), mRNA
NM_030783	Homo sapiens phosphatidylserine synthase 2 (PTDSS2), mRNA
NM_030779	Homo sapiens Eag-related gene member 2 (ERG2), mRNA
NM_030774	Homo sapiens prostate specific G-protein coupled receptor (PSGR), mRNA
NM_030772	Homo sapiens connexin 59 (GJA10), mRNA
NM_030764	Homo sapiens SH2 domain-containing phosphatase anchor protein 1 (SPAP1), mRNA
NM 030763	Homo sapiens nucleosomal binding protein 1 (NSBP1), mRNA
NM 030757	Homo sapiens makorin, ring finger protein, 4 (MKRN4), mRNA
NM_021813	Homo sapiens BTB and CNC homology 1, basic leucine zipper transcription factor 2 (BACH2), mRNA
NM 020819	Homo sapiens KIAA1411 protein (KIAA1411), mRNA
NM_030751	Homo sapiens transcription factor 8 (represses interleukin 2 expression) (TCF8), mRNA
NM 030754	Homo sapiens serum amyloid A2 (SAA2), mRNA
NM 030752	Homo sapiens t-complex 1 (TCP1), mRNA
NM 030756	Homo sapiens transcription factor 7-like 2 (T-cell specific, HMG-box)
MM_030730	(TCF7L2), mRNA
NM 006010	Homo sapiens arginine-rich, mutated in early stage tumors (ARMET), mRNA
NM 001182	Homo sapiens aldehyde dehydrogenase 7 family, member A1 (ALDH7A1),
111111_001182	mRNA
NM 000382	Homo sapiens aldehyde dehydrogenase 3 family, member A2 (ALDH3A2),
14141_000362	mRNA
NM_003486	Homo sapiens solute carrier family 7 (cationic amino acid transporter, y+
14141_003480	system), member 5 (SLC7A5), mRNA
NM_000694	Homo sapiens aldehyde dehydrogenase 3 family, member B1 (ALDH3B1),
	mRNA
NM_000693	Homo sapiens aldehyde dehydrogenase 1 family, member A3 (ALDH1A3),
	mRNA CY YO (CV VO)
NM_030381	Homo sapiens GLI-Kruppel family member GLI2 (GLI2), transcript variant 3, mRNA
	mRNA

NM_030380	Homo sapiens GLI-Kruppel family member GLI2 (GLI2), transcript variant 2, mRNA
NM_030379	Homo sapiens GLI-Kruppel family member GLI2 (GLI2), transcript variant 1, mRNA
NM_020166	Homo sapiens methylcrotonoyl-Coenzyme A carboxylase 1 (alpha) (MCCC1), mRNA
NM_005270	Homo sapiens GLI-Kruppel family member GLI2 (GLI2), transcript variant 4, mRNA
NM 002381	Homo sapiens matrilin 3 (MATN3) precursor, mRNA
NM 030583	Homo sapiens matrilin 2 (MATN2) precursor, transcript variant 2, mRNA
NM 002380	Homo sapiens matrilin 2 (MATN2) precursor, transcript variant 1, mRNA
NM 002379	Homo sapiens matrilin 1, cartilage matrix protein (MATN1), mRNA
NM_000168	Homo sapiens GLI-Kruppel family member GLI3 (Greig cephalopolysyndactyly syndrome) (GLI3), mRNA
NM 003462	Homo sapiens dynein, axonemal, light intermediate polypeptide (P28), mRNA
NM 017493	Homo seniens Hin-1 (HSHIN1), mRNA
NM_005602	Homo sapiens claudin 11 (oligodendrocyte transmembrane protein) (CLDN11), mRNA
NM_001195	Homo sapiens beaded filament structural protein 1, filensin (BFSP1), mRNA
NM 004987	Homo sapiens LIM and senescent cell antigen-like domains 1 (LIMS1), mRNA
NM 000412	Homo sapiens histidine-rich glycoprotein (HRG), mRNA
NM_024494	Homo sapiens wingless-type MMTV integration site family, member 2B (WNT2B), transcript variant WNT-2B2, mRNA
NM_004993	Homo sapiens Machado-Joseph disease (spinocerebellar ataxia 3, olivopontocerebellar ataxia 3, autosomal dominant, ataxin 3) (MJD), transcript variant 1, mRNA
NM_004185	Homo sapiens wingless-type MMTV integration site family, member 2B (WNT2B), transcript variant WNT-2B1, mRNA
NM 024415	Homo sapiens VASA protein (VASA), transcript variant 2, mRNA
NM_004398	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 10 (RNA helicase) (DDX10), mRNA
NM_004397	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 6 (RNA helicase, 54kD) (DDX6), mRNA
NM_004396	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 5 (RNA helicase, 68kD) (DDX5), mRNA
NM_030588	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 9 (RNA helicase A, nuclear DNA helicase II; leukophysin) (DDX9), transcript variant 2, mRNA
NM_001357	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 9 (RNA helicase A, nuclear DNA helicase II; leukophysin) (DDX9), transcript variant 1, mRNA
NM_004660	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide, Y chromosome (DBY), mRNA
NM_019039	Homo sapiens VASA protein (VASA), transcript variant 1, mRNA
NM_012382	Homo sapiens osmosis responsive factor (OSRF), mRNA
NM_000387	Homo sapiens solute carrier family 25 (carnitine/acylcarnitine translocase), member 20 (SLC25A20), mitochondrial protein encoded by nuclear gene, mRNA
NM_007240	Homo sapiens dual specificity phosphatase 12 (DUSP12), mRNA
NM_004940	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 7 (RNA helicase, 52kD) (DDX7), mRNA
NM 004939	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 1 (DDX1),

nRNA e 1 (MBTPS1), r, subfamily S, NA (GJA3), mRNA
e 1 (MBTPS1), r, subfamily S, NA
r, subfamily S,
NA
NA
(GJA3), mRNA
lo2, SUR4/Elo3,
NA.
ovalbumin),
NA
K2), mRNA
Α
A), mRNA
155 expressed
N. 7
N-7 protein 3
<del></del>
<i>CC:</i>
ffinity
DNIA
RNA 7, F21 and O13
, F21 and O13
, mRNA
, 11111111
mRNA
1111/1/17
etic protein 3)
no protein 3)
(MEI) mRNA
(ME1), mRNA mRNA
(ME1), mRNA , mRNA , mRNA

	mRNA CVP4F8)
NM_007253	Homo sapiens cytochrome P450, subfamily IVF, polypeptide 8 (CYP4F8),
	73.7.4
NM_000779	Homo sapiens cytochrome P450, subfamily IVB, polypeptide 1 (CYP4B1),
	mDNA
NM 001514	Homo sapiens general transcription factor IIB (GTF2B), mRNA
NM 004127	II senions G protein nathway suppressor I (GPSI), IIIXXX
NM 024423	Hame conjens desmocollin 3 (DSC3), transcript variant DSc30, IIIXIA
NM 001941	Homo copiens desmocollin 3 (DSC3), transcript variant DSc3a, mixty
NM 004949	Home copiens desmocollin 2 (DSC2), transcript variant DSc20, midva
NM 024422	Home geniens desmocollin 2 (DSC2), transcript variant Dsc2a, nikiva
NM 004948	Home gapiens desmocollin 1 (DSC1), transcript variant DSC10, IIIKNA
NM 024421	transcript Variant DSC1a, Illicia
	TT 1-mage energific DNA hinding protein 1 (12/kD) (DDD1), interior
NM_001923	TX The Tit call adhesion molecule (hydrocephalus, stellosis of aqueduot of
NM_000425	a 1 · 1 MACA (montal retardation anhasia, shiffling gail and addition
	1
> 7 C 004002	Transistant I call adhesion molecule (hydrocephalus, stellosis of aqueduct of
NM_024003	l a 1 · 1 MACA (montal retardation anhasia, shilling gait and adducted
	1 1 1 - 1 - 1 - 1 - 1 - 1 - 1 - 1 - 1 -
	Homo sapiens ferredoxin reductase (FDXR), transcript variant 2, nuclear gene
NM_004110	Homo sapiens ierredoxin reduciase (1 DAR), a amost protein mPNA
	encoding mitochondrial protein, mRNA  Homo sapiens ferredoxin reductase (FDXR), transcript variant 1, nuclear gene
NM_024417	Homo sapiens ferredoxin reductase (TDAR), utiliserile value of
	encoding mitochondrial protein, mRNA
NM_023944	Homo sapiens cytochrome P450 isoform 4F12 (CYP4F12), mRNA
NM_022845	Homo sapiens core-binding factor, beta subunit (CBFB), transcript variant 1,
	mRNA (GAN) mRNA
NM 022041	Homo sapiens giant axonal neuropathy (gigaxonin) (GAN), mRNA  Homo sapiens giant axonal neuropathy (gigaxonin) (GAN), mRNA
NM_021187	Homo sapiens cytochrome P450, subfamily IVF, polypeptide 11 (CYP4F11),
_	mRNA (TA COPI) mPNA
NM 019599	Homo sapiens taste receptor, type 2, member 1 (TAS2R1), mRNA
NM_017579	Homo sapiens taste receptor, type 2, memory 1 (DMBT1), transcript variant Homo sapiens deleted in malignant brain tumors 1 (DMBT1), transcript variant
	2 mDNA
NM 015670	Homo sapiens sentrin/SUMO-specific protease 3 (SENP3), mRNA
NM_012096	Homo sapiens adaptor protein containing pH domain, P1B domain and redeme
14141_012090	gipner motif (APPI.) mRNA
NM_005392	Tyr DIID finger protein 2 (PHF2) mRNA
NM_000896	Homo sapiens cytochrome P450, subfamily IVF, polypeptide 3 (leukotrene B4
14141_000050	amaga hydroxylase) (CYP4F3), mRNA
NM 022661	Homo sapiens SPANX family, member C (SPANXC), mRNA
NM 022573	Homo sapiens TSPYq1 (TSPYQ1), mRNA
NM 022089	Home series putative ATPase (HSA9947), mRNA
	Homo saniens hypothetical protein dJ434O14.3 (DJ434O14.3), mRNA
NM_025228	Homo sapiens KIAA1031 protein (KIAA1031), mRNA
NM_025013	Homo conjens hypothetical protein FLJ14200 (FLJ14200), IIINIA
NM_025027	TI regions hymothetical protein FI 120958 (FLJ20958), MKNA
NM_022102	to the manufacture of the manufa
NM_021724	
	mRNA Homo sapiens hypothetical protein MGC10902 (MGC10902), mRNA
NM_030570	
NM_025135	
NM_024317	
NM 021822	Homo sapiens phorbolin-like protein WD3013 (WD3013), interin

ND 6 017600	II A CO for paris a material homologic (HCDNA CDH) mDNA
NM_017509	Homo sapiens ACO for serine protease homologue (HSRNASPH), mRNA
NM_005583	Homo sapiens lymphoblastic leukemia derived sequence 1 (LYL1), mRNA
NM_020070	Homo sapiens immunoglobulin lambda-like polypeptide 1 (IGLL1), mRNA
NM_002383	Homo sapiens MYC-associated zinc finger protein (purine-binding transcription
ND4 016044	factor) (MAZ), mRNA
NM_016944	Homo sapiens taste receptor, type 2, member 4 (TAS2R4), mRNA
NM_016943	Homo sapiens taste receptor, type 2, member 3 (TAS2R3), mRNA
NM_000378	Homo sapiens Wilms tumor 1 (WT1), transcript variant A, mRNA
NM_024426	Homo sapiens Wilms tumor 1 (WT1), transcript variant D, mRNA
NM_024425	Homo sapiens Wilms tumor 1 (WT1), transcript variant C, mRNA
NM_024424	Homo sapiens Wilms tumor 1 (WT1), transcript variant B, mRNA
NM_000765	Homo sapiens cytochrome P450, subfamily IIIA, polypeptide 7 (CYP3A7), mRNA
NM_021570	Homo sapiens BarH-like homeobox 1 (BARX1), mRNA
NM_000068	Homo sapiens calcium channel, voltage-dependent, P/Q type, alpha 1A subunit
_	(CACNA1A), transcript variant 1, mRNA
NM_030574	Homo sapiens hypothetical protein MGC10327 (MGC10327), mRNA
NM_030573	Homo sapiens hypothetical protein MGC10963 (MGC10963), mRNA
NM_024867	Homo sapiens hypothetical protein FLJ23577 (FLJ23577), mRNA
NM_002739	Homo sapiens protein kinase C, gamma (PRKCG), mRNA
NM_020548	Homo sapiens diazepam binding inhibitor (GABA receptor modulator, acyl-
	Coenzyme A binding protein) (DBI), mRNA
NM_025176	Homo sapiens KIAA0980 protein (KIAA0980), mRNA
NM_003789	Homo sapiens TNFRSF1A-associated via death domain (TRADD), mRNA
NM_017541	Homo sapiens crystallin, gamma S (CRYGS), mRNA
NM_006891_	Homo sapiens crystallin, gamma D (CRYGD), mRNA
NM_020989	Homo sapiens crystallin, gamma C (CRYGC), mRNA
NM_005210	Homo sapiens crystallin, gamma B (CRYGB), mRNA
NM_014617	Homo sapiens crystallin, gamma A (CRYGA), mRNA
NM_002396	Homo sapiens malic enzyme 2, NAD(+)-dependent, mitochondrial (ME2),
	nuclear gene encoding mitochondrial protein, mRNA
NM_025268	Homo sapiens hypothetical protein MGC4659 (MGC4659), mRNA
NM_025244	Homo sapiens testis specific, 10 (TSGA10), mRNA
NM_025240	Homo sapiens B7 homolog 3 (B7-H3), mRNA
NM_025237	Homo sapiens sclerostin (SOST), mRNA
NM_025236	Homo sapiens HZFw1 protein (HZFW1), mRNA
NM_025235	Homo sapiens tankyrase 2 (TNKL), mRNA
NM_025233	Homo sapiens nucleotide binding protein (NBP), mRNA
NM_025232	Homo sapiens hypothetical protein FLJ22246 (FLJ22246), mRNA
NM_025218	Homo sapiens UL16-binding protein 1 (ULBP1), mRNA
NM_025217	Homo sapiens UL16-binding protein 2 (ULBP2), mRNA
NM_025215	Homo sapiens pseudouridine synthase 1 (PUS1), mRNA
NM_025214	Homo sapiens CTCL tumor antigen se57-1 (SE57-1), mRNA
NM_025212	Homo sapiens Dvl-binding protein IDAX (inhibition of the Dvl and Axin
	complex) (IDAX), mRNA
NM_025210	Homo sapiens type 1 protein phosphatase inhibitor (I-4), mRNA
NM_025209	Homo sapiens enhancer of polycomb 1 (EPC1), mRNA
NM_025205	Homo sapiens hypothetical protein DKFZp434N185 (DKFZP434N185), mRNA
NM_025198	Homo sapiens transcription termination factor-like protein (LOC80298), mRNA
NM_025193	Homo sapiens 3 beta-hydroxy-delta 5-C27-steroid oxidoreductase (C(27)-3BETA-HSD), mRNA
NM 025180	Homo sapiens hypothetical protein FLJ13386 (FLJ13386), mRNA
	The state of the s

	77 7001775 (EL 100175) mDNA
NM 025161	Homo sapiens hypothetical protein FLJ22175 (FLJ22175), mRNA
NM 025158	II ama conjens hymothetical protein FLJ22251 (FLJ22251), IIICIVI
NM 025148	Homo sapiens hypothetical protein FLJ12986 (FLJ12986), mRNA Homo sapiens hypothetical protein FLJ12986 (FLJ12986), mRNA
NM 025137	Homo sapiens hypothetical protein FLJ21439 (FLJ21439), mRNA Homo sapiens hypothetical protein FLJ21439 (FLJ21439), mRNA
NM 025116	Times conjens hymothetical protein FLJ12/81 (FLJ12/81), IIIXIA
NM 025114	Homo sapiens hypothetical protein FLJ13615 (FLJ13615), mRNA Homo sapiens hypothetical protein FLJ13615 (FLJ13815), mRNA
NM 025083	Homo seniens hypothetical protein FLJ21128 (FLJ21128), HKVA
NM 025054	Homo sapiens hypothetical protein FLJ23132 (FLJ23132), mRNA
NM 025017	Homo sapiens hypothetical protein FLJ13892 (FLJ13892), mRNA  Homo sapiens hypothetical protein FLJ13892 (FLJ13892), mRNA
NM 025011	Homo caniens hypothetical protein FLJ13/44 (FLJ13/44), illicity
NM 024995	Homo sapiens hypothetical protein FLJ12616 (FLJ12616), mRNA
NM 024987	Homo sapiens hypothetical protein FLJ12345 (FLJ12345), mRNA  Homo sapiens hypothetical protein FLJ12345 (FLJ12345), mRNA
NM 024900	Homo sapiens hypothetical protein FLJ22479 (FLJ22479), mRNA  Homo sapiens hypothetical protein FLJ22479 (FLJ22479), mRNA
NM 024874	Homo sapiens hypothetical protein FLJ14225 (FLJ14225), mRNA  Homo sapiens hypothetical protein FLJ14225 (FLJ14225), mRNA
NM 024873	Homo sapiens hypothetical protein FLJ21162 (FLJ21162), mRNA  Homo sapiens hypothetical protein FLJ21162 (FLJ21162), mRNA
NM 024861	Homo sapiens hypothetical protein FLJ22671 (FLJ22671), mRNA  Homo sapiens hypothetical protein FLJ22671 (FLJ22671), mRNA
NM 024836	Homo sapiens hypothetical protein FLJ22301 (FLJ22301), mRNA  Homo sapiens hypothetical protein FLJ22301 (FLJ22301), mRNA
NM 024822	Homo sapiens hypothetical protein FLJ22601 (FLJ22601), mRNA  Homo sapiens hypothetical protein FLJ22601 (FLJ22651), mRNA
NM 024819	Homo sapiens hypothetical protein FLJ22955 (FLJ22955), mRNA  Homo sapiens hypothetical protein FLJ22955 (FLJ22955), mRNA
NM 024816	Homo sapiens hypothetical protein FLJ23282 (FLJ23282), mRNA  Homo sapiens hypothetical protein FLJ23282 (FLJ23282), mRNA
NM 024803	Homo sapiens hypothetical protein FLJ21665 (FLJ21665), mRNA
NM 024795	Homo sapiens hypothetical protein FLJ22800 (FLJ22800), mRNA
NM 024767	Homo sapiens hypothetical protein FLJ21120 (FLJ21120), mRNA  Homo sapiens hypothetical protein FLJ21120 (FLJ21120), mRNA
NM_024760	Homo sapiens hypothetical protein FLJ14009 (FLJ14009), mRNA  Homo sapiens hypothetical protein FLJ14009 (FLJ14009), mRNA
NM_024741	Homo sapiens hypothetical protein FLJ12827 (FLJ12827), mRNA  Homo sapiens hypothetical protein FLJ12827 (FLJ12827), mRNA
NM_024723	Homo sapiens hypothetical protein FLJ23471 (FLJ23471), mRNA  Homo sapiens hypothetical protein FLJ23471 (FLJ23471), mRNA
NM_024720	Homo sapiens hypothetical protein FLJ23510 (FLJ23510), mRNA  Homo sapiens hypothetical protein FLJ23510 (FLJ23510), mRNA
NM_024698	Homo sapiens hypothetical protein FLJ13044 (FLJ13044), mRNA  Homo sapiens hypothetical protein FLJ13044 (FLJ13049), mRNA
NM_024692	Homo sapiens hypothetical protein FLJ21069 (FLJ21069), mRNA  Homo sapiens hypothetical protein FLJ21069 (FLJ21069), mRNA
NM_024689	Homo sapiens hypothetical protein FLJ14103 (FLJ14103), mRNA  Homo sapiens hypothetical protein FLJ14103 (FLJ14103), mRNA
NM_024687	Homo sapiens hypothetical protein FLJ23049 (FLJ23049), mRNA  Homo sapiens hypothetical protein FLJ23049 (FLJ23049), mRNA
NM_024648	Homo sapiens hypothetical protein FLJ22222 (FLJ22222), mRNA  Homo sapiens hypothetical protein FLJ22222 (FLJ22222), mRNA
NM_024622	Homo sapiens hypothetical protein FLJ21901 (FLJ21901), mRNA  Homo sapiens hypothetical protein FLJ21901 (FLJ21901), mRNA
NM_024611	Homo sapiens hypothetical protein FLJ11896 (FLJ11896), mRNA  Homo sapiens hypothetical protein FLJ11896 (FLJ11896), mRNA
NM_024591	Homo sapiens hypothetical protein FLJ11749 (FLJ11749), mRNA  Homo sapiens hypothetical protein FLJ12054 (FLJ2054) mRNA
NM_024561	Homo sapiens hypothetical protein FLJ22054 (FLJ22054), mRNA  Homo sapiens hypothetical protein FLJ22054 (FLJ22054), mRNA
NM_024540	Homo sapiens hypothetical protein FLJ20917 (FLJ20917), mRNA  Homo sapiens hypothetical protein FLJ20917 (FLJ20917), mRNA
NM_024518	Homo sapiens UL16-binding protein 3 (ULBP3), mRNA  Homo sapiens UL16-binding protein MCC4645 (MCC4645) mRNA
NM_024515	Homo sapiens hypothetical protein MGC4645 (MGC4645), mRNA  Homo sapiens hypothetical protein mGC4645 (MGC4645), mRNA
NM_024504	Homo sapiens PR domain containing 14 (PRDM14), mRNA
NM_024501	Homo sapiens homeo box D1 (HOXD1), mRNA  Homo sapiens peroxisomal long-chain acyl-coA thioesterase (ZAP128), mRNA
NM_006821	
NM_006680	
	mRNA Homo sapiens desmoglein 3 (pemphigus vulgaris antigen) (DSG3), mRNA
NM_001944	
NM_001943	
NM_001942	
NM_024500	
NM_024498	
NM_018943	
NM_015640	nomo sapiens rai-i inclari-oniumg protein (122 20)

NM_015332 NM_022001 NM_021708	Homo sapiens KIAA1068 protein (KIAA1068), mRNA  Homo sapiens SMAD in the antisense orientation (DAMS), mRNA
	HOMO Sablens NVIALL in the anticence orientation (LIAMX) mVNIA
NM_021708	
	Homo sapiens leukocyte-associated Ig-like receptor 1 (LAIR1), transcript variant d, mRNA
NM_021706	Homo sapiens leukocyte-associated Ig-like receptor 1 (LAIR1), transcript variant b, mRNA
NM_002287	Homo sapiens leukocyte-associated Ig-like receptor 1 (LAIR1), transcript variant a, mRNA
NM_004424	Homo sapiens E4F transcription factor 1 (E4F1), mRNA
NM 018834	Homo sapiens matrin 3 (MATR3), mRNA
NM_017830	Homo sapiens ovarian carcinoma immunoreactive antigen (OCIA), mRNA
NM 006926	Homo sapiens surfactant, pulmonary-associated protein A2 (SFTPA2), mRNA
NM 005411	Homo sapiens surfactant, pulmonary-associated protein A1 (SFTPA1), mRNA
NM 024492	Homo sapiens apolipoprotein (a) related gene C (APOARGC), mRNA
NM 024491	Homo sapiens p10-binding protein (BITE), mRNA
NM_015472	Homo sapiens transcriptional co-activator with PDZ-binding motif (TAZ) (TAZ), mRNA
NM 017797	Homo sapiens BTB (POZ) domain containing 2 (BTBD2), mRNA
NM_002826	Homo sapiens quiescin Q6 (QSCN6), mRNA
NM_024010	Homo sapiens 5-methyltetrahydrofolate-homocysteine methyltransferase
_	reductase (MTRR), transcript variant 2, mRNA
NM_004972	Homo sapiens Janus kinase 2 (a protein tyrosine kinase) (JAK2), mRNA
NM_000761	Homo sapiens cytochrome P450, subfamily I (aromatic compound-inducible), polypeptide 2 (CYP1A2), mRNA
NM_000104	Homo sapiens cytochrome P450, subfamily I (dioxin-inducible), polypeptide 1 (glaucoma 3, primary infantile) (CYP1B1), mRNA
NM_000499	Homo sapiens cytochrome P450, subfamily I (aromatic compound-inducible), polypeptide 1 (CYP1A1), mRNA
NM 024318	Homo sapiens immunoglobulin-like transcript 8 (ILT8), mRNA
NM 021806	Homo sapiens 2.19 gene (2.19), mRNA
NM_006208	Homo sapiens ectonucleotide pyrophosphatase/phosphodiesterase 1 (ENPP1), mRNA
NM_007076	Homo sapiens Huntingtin interacting protein E (HYPE), mRNA
NM_018571	Homo sapiens amyotrophic lateral sclerosis 2 (juvenile) chromosome region, candidate 2 (ALS2CR2), mRNA
NM_015049	Homo sapiens amyotrophic lateral sclerosis 2 (juvenile) chromosome region, candidate 3 (ALS2CR3), mRNA
NM 023036	Homo sapiens dynein intermediate chain 2 (DNAI2), mRNA
NM 022171	Homo sapiens T-cell leukemia translocation altered gene (TCTA), mRNA
NM 016128	Homo sapiens coat protein gamma-cop (LOC51137), mRNA
NM 021999	Homo sapiens integral membrane protein 2B (ITM2B), mRNA
NM_021992	Homo sapiens thymosin, beta, identified in neuroblastoma cells (TMSNB), mRNA
NM 021994	Homo sapiens zinc finger protein 277 (ZNF277), mRNA
NM 007257	Homo sapiens paraneoplastic antigen MA2 (PNMA2), mRNA
NM 021972	Homo sapiens sphingosine kinase 1 (SPHK1), mRNA
NM_021976	Homo sapiens retinoid X receptor, beta (RXRB), mRNA
NM 021963	Homo sapiens nucleosome assembly protein 1-like 2 (NAP1L2), mRNA
NM_021978	Homo sapiens suppression of tumorigenicity 14 (colon carcinoma, matriptase, epithin) (ST14), mRNA
NM_021977	Homo sapiens solute carrier family 22 (extraneuronal monoamine transporter), member 3 (SLC22A3), mRNA

	140 ( HZ 52) (ZNE148) mRNA
VM 021964 1	Homo sapiens zinc finger protein 148 (pHZ-52) (ZNF148), mRNA
VM 021966	Homo sapiens T-cell leukemia/lymphoma 1A (TCL1A), mRNA
NM 012186	Homo sapiens forkhead box E3 (FOXE3), mRNA
NM 012182	Homo sapiens forkhead box B1 (FOXB1), mRNA
	II a serious ligatin (I GTN) mRNA
TO 6 001055	Homo saniens guanine nucleotide binding protein (G protein), gainina
_	
NM_021959	Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 11
	(DDD1D11) DNIA
NM_021951	Homo sapiens doublesex and mab-3 related transcription factor 1 (DMRT1),
	7 A T A
NM_021960	mRNA  Homo sapiens myeloid cell leukemia sequence 1 (BCL2-related) (MCL1),
14147_021900	73.T A
NM_021952	Homo sapiens ELAV (embryonic lethal, abnormal vision, Drosophila)-like 4 (Hu
14141_021552	
NM_021949	Homo sapiens ATPase, Ca++ transporting, plasma membrane 3 (ATP2B3),
14141_021545	mRNA
NM 021953	IV comicons for liberal box M1 (FOXM1), mRNA
NM 021956	in the most recentor ionotropic kainate 2 (UKIK2), IIIKNA
NM 004886	Homo sapiens amyloid beta (A4) precursor protein-binding, family A, member 3
14141_004666	$\alpha z_{11} + \alpha z_{12} = \alpha z_{13} + \alpha z_{13} + \alpha z_{13} = \alpha z_{13} + \alpha z_{13} = \alpha z_{13} + \alpha z_{13} = \alpha z_{13} $
NM_006557	Homo sapiens doublesex and mab-3 related transcription factor 2 (DMRT2),
MMT_000221	DATA
ND ( 000052	Homo sapiens kinase insert domain receptor (a type III receptor tyrosine kinase)
NM_002253	(KEDD)DNIA
ND 4 000179	ry in a inquire like growth factor hinding protein 6 (IGFBPO), IIINNA
NM_002178	Homo sapiens succinate-CoA ligase, ADP-forming, beta subunit (SUCLA2),
NM_003850	I ====:
NT ( 002802	Home saniers myosin heavy polypeptide 13, skeletal muscle (MYH13), mKNA
NM_003802	Homo sapiens zinc finger protein 16 (KOX 9) (ZNF16), mRNA
NM_006958	Homo sapiens tousled-like kinase 2 (TLK2), mRNA
NM_006852	Homo sapiens netrin 4 (NTN4), mRNA
NM_021229	Homo sapiens NADPH oxidase 3 (NOX3), mRNA
NM_015718	Homo sapiens NADI II Oxidase 5 (NOA5), MECH
NM_015003	Homo sapiens golgin-67 (KIAA0855), mRNA  Homo sapiens N-ethylmaleimide-sensitive factor (NSF), mRNA
NM_006178	Homo sapiens N-ethylmaleimide-sensitive factor (1457), and the lattice of (
NM_003116	Homo sapiens sperm associated antigen 4 (SPAG4), mRNA
NM_018724	Homo sapiens interleukin 20 (IL20), mRNA
NM_019083	Homo sapiens hypothetical protein (FLJ10287), mRNA
NM 003114	Homo sapiens sperm associated antigen 1 (SPAG1), mRNA  Homo sapiens sperm associated antigen 1 (SPAG1), member 1
NM 021097	Homo sapiens solute carrier family 8 (sodium/calcium exchanger), member 1
_	(SLC8A1), mRNA
NM 021102	Homo sapiens serine protease inhibitor, Kunitz type, 2 (SPINT2), mRNA
NM 021101	TI was assigned algorithm 1 (CLDN1) mRNA
NM 021095	Homo sapiens claudin 1 (CEDIVI), index 1  Homo sapiens solute carrier family 5 (sodium-dependent vitamin transporter),
	1 6 (SI C5 λ6) mPNΔ
NM_021076	Homo saniens neurofilament, heavy polypeptide (200kD) (NEFH), mRNA
NM 001751	TTiona aveteinul tRNA synthetase (CAKS), IIINNA
1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	Homo sapiens VADH dehydrogenase (ubiquinone) flavoprotein 2 (24kD)
	The same of the sa
NM_021074	(NDUFV2), mRNA
NM_021074	(NDUFV2), mRNA  Homo sapiens macrophage stimulating 1 (hepatocyte growth factor-like)
	(NDUFV2), mRNA  Homo sapiens macrophage stimulating 1 (hepatocyte growth factor-like)  (MST1), mRNA  Homo sapiens synovial sarcoma, X breakpoint 2 (SSX2), mRNA

NM_015392	Homo sapiens neural proliferation, differentiation and control, 1 (NPDC1), mRNA
NM 020482	Homo sapiens activator of CREM in testis (ACT), mRNA
NM 014509	Homo sapiens kraken-like (BK126B4.1), mRNA
NM_005132	Homo sapiens Rec8p, a meiotic recombination and sister chromatid cohesion phosphoprotein of the rad21p family (REC8), mRNA
NM_018896	Homo sapiens calcium channel, voltage-dependent, alpha 1G subunit (CACNA1G), mRNA
NM_005329	Homo sapiens hyaluronan synthase 3 (HAS3), mRNA
NM_015193	Homo sapiens activity-regulated cytoskeleton-associated protein (ARC), mRNA
NM_016203	Homo sapiens protein kinase, AMP-activated, gamma 2 non-catalytic subunit (PRKAG2), mRNA
NM_000627	Homo sapiens latent transforming growth factor beta binding protein 1 (LTBP1), mRNA
NM_002454	Homo sapiens 5-methyltetrahydrofolate-homocysteine methyltransferase reductase (MTRR), transcript variant 1, mRNA
NM_001091	Homo sapiens amiloride binding protein 1 (amine oxidase (copper-containing)) (ABP1), mRNA
NM 024016	Homo sapiens homeo box B8 (HOXB8), mRNA
NM 024015	Homo sapiens homeo box B4 (HOXB4), mRNA
NM 015227	Homo sapiens KIAA0958 protein (KIAA0958), mRNA
NM_024430	Homo sapiens proline-serine-threonine phosphatase interacting protein 2 (PSTPIP2), mRNA
NM 003588	Homo sapiens cullin 4B (CUL4B), mRNA
NM 016059	Homo sapiens peptidylprolyl isomerase (cyclophilin)-like 1 (PPIL1), mRNA
NM 014432	Homo sapiens interleukin 20 receptor, alpha (IL20RA), mRNA
NM 000270	Homo sapiens nucleoside phosphorylase (NP), mRNA
NM_003021	Homo sapiens small glutamine-rich tetratricopeptide repeat (TPR)-containing (SGT), mRNA
NM_002038	Homo sapiens interferon, alpha-inducible protein (clone IFI-6-16) (G1P3), transcript variant 1, mRNA
NM_022873	Homo sapiens interferon, alpha-inducible protein (clone IFI-6-16) (G1P3), transcript variant 3, mRNA
NM_022872	Homo sapiens interferon, alpha-inducible protein (clone IFI-6-16) (G1P3), transcript variant 2, mRNA
NM_022803	Homo sapiens uncoupling protein 3 (mitochondrial, proton carrier) (UCP3), transcript variant short, nuclear gene encoding mitochondrial protein, mRNA
NM_003356	Homo sapiens uncoupling protein 3 (mitochondrial, proton carrier) (UCP3), transcript variant long, nuclear gene encoding mitochondrial protein, mRNA
NM_022810	Homo sapiens solute carrier family 25 (mitochondrial carrier, brain), member 14 (SLC25A14), transcript variant short, nuclear gene encoding mitochondrial protein, mRNA
NM_003355	Homo sapiens uncoupling protein 2 (mitochondrial, proton carrier) (UCP2), nuclear gene encoding mitochondrial protein, mRNA
NM_021833	Homo sapiens uncoupling protein 1 (mitochondrial, proton carrier) (UCP1), nuclear gene encoding mitochondrial protein, mRNA
NM_002231	Homo sapiens kangai 1 (suppression of tumorigenicity 6, prostate; CD82 antigen (R2 leukocyte antigen, antigen detected by monoclonal and antibody IA4)) (KAI1), mRNA
NM_004967	Homo sapiens integrin-binding sialoprotein (bone sialoprotein, bone sialoprotein II) (IBSP), mRNA
NM 000490	Homo sapiens arginine vasopressin (neurophysin II, antidiuretic hormone,

	diabetes insipidus, neurohypophyseal) (AVP), mRNA
VM_022877	Homo sapiens survival of motor neuron 2, centrollier (SM142), dancer.
_	· · · · · · · · · · · · · · · · · · ·
VM 022876	Homo sapiens survival of motor neuron 2, centromeric (SMN2), transcript
	· 11DXIA
VM 022875	Homo sapiens survival of motor neuron 2, centromeric (SMN2), transcript
	· · Daia
NM 017411	Homo sapiens survival of motor neuron 2, centromeric (SMN2), transcript
1111_027	variant d. mRNA
NM 005474	Homo saniens histone deacetylase 5 (HDAC5), mRNA
NM 006037	history descentilese 4 (HDAC4), mKNA
NM_003474	Homo sapiens a disintegrin and metalloproteinase domain 12 (meltrin alpha)
14147_002414	Livering sand in the second of 1 mVNA
NM_000344	Homo sapiens survival of motor neuron 1, telomeric (SMN1), transcript variant
IAIMT_00024-4	1 4 95374
ND 6 022974	d, mRNA  Homo sapiens survival of motor neuron 1, telomeric (SMN1), transcript variant
NM_022874	b, mRNA
27.6.006400	1 demostin 2 (n50) (DCTN2) mRNA
NM_006400	Homo sapiens nuclear receptor subfamily 0, group B, member 2 (NR0B2),
NM_021969	DAIA
	mRNA Homo sapiens small EDRK-rich factor 1A (telomeric) (SERF1A), mRNA
NM_021967	Homo sapiens general transcription factor IIH, polypeptide 2 (44kD subunit)
NM_001515	Homo sapiens general transcription factor into polyperate
	(GTF2H2), mRNA  Homo sapiens solute carrier family 25 (mitochondrial carrier, brain), member 14
NM_003951	Homo sapiens solute carrier family 25 (introchondrial carrier, standy,
	(SLC25A14), transcript variant long, nuclear gene choosing missions
	protein, mRNA
NM_004277	Homo sapiens uncoupling protein 4 (UCP4), nuclear gene encoding
	mitochondrial protein, mRNA
NM 004536	Homo sapiens baculoviral IAP repeat-containing 1 (BIRC1), mRNA
NM_000346	Homo sapiens SRY (sex determining region Y)-box 9 (camponicite dysplasia,
_	1 morromol) (S() X 9) mR NA
NM_003645	Homo sapiens fatty-acid-Coenzyme A ligase, very long-chain 1 (FACVL1),
	mPNA
NM 024409	Homo sapiens natriuretic peptide precursor C (NPPC), mRNA
NM 024410	Try assigns outer dense fibre of sperm fails I (ODFI), MIKINA
NM 004180	Homo sapiens TRAF family member-associated NFKB activator (TANK),
14141_00+100	mRNA
NM_024332	Hame genions of 1A (C6 1A) mRNA
NM_024332	Homo seniens hypothetical protein MGC11256 (MGC11256), HIKNA
	Homo sapiens hypothetical protein MGC4175 (MGC4175), mRNA
NM_024315	Homo seniens hypothetical protein ET (ET), mRNA
NM_024311	Homo sapiens hypothetical protein MGC4289 (MGC4289), mRNA
NM_024309	
NM_024306	Homo sapiens hypothetical protein MGC2217 (MGC2217), mRNA
NM_024300	
NM_024296	
NM_024294	
NM_024292	
NM_024012	T C
NM_024123	Homo sapiens putative Ly-6 superfamily member (GoE), mixva
NM_021904	Homo sapiens gamma-aminobutyric acid (GABA) B receptor, I (GABBRI),
	DNA
NM 021903	Homo sapiens gamma-aminobutyric acid (GABA) B receptor, 1 (GABBR1),

NM_001470	transcript variant 2, mRNA
14141_001470	Homo sapiens gamma-aminobutyric acid (GABA) B receptor, 1 (GABBR1),
NM_001858	transcript variant 1, mRNA
NM 015071	Homo sapiens collagen, type XIX, alpha 1 (COL19A1), mRNA
14141_0130/1	Homo sapiens GTPase regulator associated with the focal adhesion kinase
NM 007329	_   pp123(FAK); KIAA0621 protein (KIAA0621) mRNA
11111_00/329	Homo sapiens deleted in malignant brain tumors 1 (DMBT1), transcript variant
NM_023004	z, mkiva
NM 005371	Homo sapiens nogo receptor (NOGOR), mRNA
NM 023033	Homo sapiens methyltransferase-like 1 (METTL1), transcript variant 1, mRNA
NM_023032	Fromo sapiens methyltransferase-like 1 (METTL1), transcript variant 3, mPNA
NM 014289	riomo sapiens methyltransferase-like 1 (METTL1), transcript variant 2 mRNA
NM 023089	Homo sapiens calpain 6 (CAPN6), mRNA
NM 023089	Homo sapiens calpain 10 (CAPN10), transcript variant 7, mRNA
NM_023087	Homo sapiens calpain 10 (CAPN10), transcript variant 6, mRNA
	Homo sapiens calpain 10 (CAPN10), transcript variant 5, mRNA
NM_023086 NM_023085	Homo sapiens calpain 10 (CAPN10), transcript variant 4, mRNA
	Homo sapiens calpain 10 (CAPN10), transcript variant 3, mRNA
NM 023084	Homo sapiens calpain 10 (CAPN10), transcript variant 2, mRNA
NM 023083	Homo sapiens calpain 10 (CAPN10), transcript variant 1, mRNA
NM_021251	Homo sapiens calpain 10 (CAPN10), transcript variant 8 mRNA
NM_005083	Homo sapiens U2 small nuclear ribonucleoprotein auxiliary factor, small subunit
NM_023031	1 (UZAFIRSI), mRNA
14141_023031	Homo sapiens fibroblast growth factor receptor 2 (bacteria-expressed kinase,
	Relatified the growth factor receptor, craniofacial dysostosis 1. Crouzon
	syndrome, Pfeiffer syndrome, Jackson-Weiss syndrome) (FGFR2), transcript variant 13, mRNA
NM_023030	Homo saniens fibroblest growth 6 - 4 - 2 G
= 12.12_52250	Homo sapiens fibroblast growth factor receptor 2 (bacteria-expressed kinase,
	keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon syndrome, Pfeiffer syndrome, Jackson-Weiss syndrome) (FGFR2), transcript
	variant 12, mRNA
NM_023028	Homo sapiens fibroblast growth factor receptor 2 (bacteria-expressed kinase,
_	keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon
	syndrome, Pfeiffer syndrome, Jackson-Weiss syndrome) (FGFR2), transcript
	variant 10, mRNA
NM_022976	Homo sapiens fibroblast growth factor receptor 2 (bacteria-expressed kinase,
	keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon
ı	syndrome, Pfeiffer syndrome, Jackson-Weiss syndrome) (FGFR2), transcript
	variant 9, mr.nA
NM_022975	Homo sapiens fibroblast growth factor receptor 2 (bacteria-expressed kinase,
	keramocyte growth factor receptor, cramofacial dysostosis 1. Crouzon
	syndrome, Pieilier syndrome, Jackson-Weiss syndrome) (FGFR2) transcript
ND 6 00000	variant 8, mrina
NM_022974	Homo sapiens fibroblast growth factor receptor 2 (bacteria-expressed kinase,
	Relatinocyte growin factor receptor, craniofacial dysostosis 1 Crouzon
	syndrome, Pieiffer syndrome, Jackson-Weiss syndrome) (FGFR2) transcript
NIM 022272	variant 7, mknA
NM_022973	Homo sapiens fibroblast growth factor receptor 2 (bacteria-expressed kinase,
İ	Relatifice the growth factor receptor, craniofacial dysostosis 1. Crougon
ĺ	syndrome, Pieirier syndrome, Jackson-Weiss syndrome) (FGFR2) transcript
NM_022972	variant o, mriva
1111 022912	Homo sapiens fibroblast growth factor receptor 2 (bacteria-expressed kinase,

	keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon syndrome, Pfeiffer syndrome, Jackson-Weiss syndrome) (FGFR2), transcript
	variant 5 mRNA
NM_022971	Homo sapiens fibroblast growth factor receptor 2 (bacteria-expressed kinase,
11112_0	keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon
•	syndrome, Pfeiffer syndrome, Jackson-Weiss syndrome) (FGFR2), transcript
	variant 4 mRNA
NM 022970	Homo sapiens fibroblast growth factor receptor 2 (bacteria-expressed kinase,
14141_022970	keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon
	syndrome, Pfeiffer syndrome, Jackson-Weiss syndrome) (FGFR2), transcript
	variant 3, mRNA
> Tr 6 000060	Homo sapiens fibroblast growth factor receptor 2 (bacteria-expressed kinase,
NM_022969	keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon
	syndrome, Pfeiffer syndrome, Jackson-Weiss syndrome) (FGFR2), transcript
	variant 2, mRNA
NM_015850	Homo sapiens fibroblast growth factor receptor 1 (fms-related tyrosine kinase 2,
	Pfeiffer syndrome) (FGFR1), transcript variant 2, mRNA
NM 023111	Homo sapiens fibroblast growth factor receptor 1 (fms-related tyrosine kinase 2,
	Pfeiffer syndrome) (FGFR1), transcript variant 9, mRNA
NM_023110	Homo sapiens fibroblast growth factor receptor 1 (fms-related tyrosine kinase 2,
	Pfeiffer syndrome) (FGFR1), transcript variant 8, mRNA
NM_023109	Homo sapiens fibroblast growth factor receptor 1 (fms-related tyrosine kinase 2,
14141_025105	Pfeiffer syndrome) (FGFR1), transcript variant 7, mRNA
NM 023029	Homo sapiens fibroblast growth factor receptor 2 (bacteria-expressed kinase,
14101_023029	keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon
	syndrome, Pfeiffer syndrome, Jackson-Weiss syndrome) (FGFR2), transcript
	variant 11, mRNA
27.4 022108	Homo sapiens fibroblast growth factor receptor 1 (fms-related tyrosine kinase 2,
NM_023108	Pfeiffer syndrome) (FGFR1), transcript variant 6, mRNA
	Pietier syndrome) (PGFRI), transcript variant 0, intervi
NM_000141	Homo sapiens fibroblast growth factor receptor 2 (bacteria-expressed kinase,
	keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon
	syndrome, Pfeiffer syndrome, Jackson-Weiss syndrome) (FGFR2), transcript
	variant 1, mRNA
NM_023107	Homo sapiens fibroblast growth factor receptor 1 (fms-related tyrosine kinase 2,
	Pfeiffer syndrome) (FGFR1), transcript variant 5, mRNA
NM 023106	Homo sapiens fibroblast growth factor receptor 1 (fms-related tyrosine kinase 2,
	Pfeiffer syndrome) (FGFR1), transcript variant 4, mRNA
NM 023105	Homo sapiens fibroblast growth factor receptor 1 (fms-related tyrosine kinase 2,
_	Pfeiffer syndrome) (FGFR1), transcript variant 3, mRNA
NM_000604	Homo sapiens fibroblast growth factor receptor 1 (fms-related tyrosine kinase 2,
1111_00000.	Pfeiffer syndrome) (FGFR1), transcript variant 1, mRNA
NM 024018	Homo sapiens butyrophilin, subfamily 2, member A3 (BTN2A3), mRNA
	Homo sapiens betaine-homocysteine methyltransferase 2 (BHMT2), mRNA
NM_017614	Homo sapiens BENE protein (BENE), mRNA
NM_005434	Homo sapiens steroid sulfatase (microsomal), arylsulfatase C, isozyme S (STS),
NM_000351	mRNA
NM 024105	Homo sapiens hypothetical protein MGC3136 (MGC3136), mRNA
	Homo sapiens hypothetical protein MGC2574 (MGC2574), mRNA
NM_024098	Homo sapiens hypothetical protein MGC5627 (MGC5627), mRNA
NM_024096	Homo sapiens nypoinetical protein MCC5540 (MCC5540), mRNA
NM_024095	Homo sapiens hypothetical protein MGC5540 (MGC5540), mRNA
NM_024091	Homo sapiens hypothetical protein MGC5297 (MGC5297), mRNA
NM_024089	Homo sapiens hypothetical protein MGC5302 (MGC5302), mRNA

NM 024082	Homo sapiens transmembrane gamma-carboxyglutamic acid protein 3 (TMG3),
	mRNA
NM_024081	Homo sapiens transmembrane gamma-carboxyglutamic acid protein 4 (TMG4), mRNA
NM_024079	Homo sapiens hypothetical protein MGC2840 similar to a putative glucosyltransferase (MGC2840), mRNA
NM_024078	Homo sapiens hypothetical protein MGC3162 (MGC3162), mRNA
NM_024075	Homo sapiens LENG5 protein (LENG5), mRNA
NM_024073	Homo sapiens hypothetical protein MGC2875 (MGC2875), mRNA
NM 024060	Homo sapiens hypothetical protein MGC5395 (MGC5395), mRNA
NM 024056	Homo sapiens hypothetical protein MGC5576 (MGC5576), mRNA
NM 024054	Homo sapiens hypothetical protein MGC2821 (MGC2821), mRNA
NM 024051	Homo sapiens hypothetical protein MGC3077 (MGC3077), mRNA
NM 024047	Homo sapiens hypothetical protein MGC3037 (MGC3037), mRNA
NM 024044	Homo sapiens hypothetical protein MGC5178 (MGC5037), mRNA
NM 024043	Homo sapiens hypothetical protein MGC3101 (MGC3101), mRNA
NM 024035	Homo sapiens hypothetical protein MGC3113 (MGC3113), mRNA
NM 024034	Homo sapiens hypothetical protein MGC3129 similar to ganglioside-induced
	differentiation-associated protein (MGC3129), mRNA
NM 024009	Homo sapiens gap junction protein, beta 3, 31kD (connexin 31) (GJB3), mRNA
NM 024013	Homo sapiens interferon, alpha 1 (IFNA1), mRNA
NM 000521	Homo sapiens hexosaminidase B (beta polypeptide) (HEXB), mRNA
NM 000520	Homo sapiens hexosaminidase A (alpha polypeptide) (HEXA), mRNA
NM 006044	Homo sapiens histone deacetylase 6 (HDAC6), mRNA
NM 003883	Homo sapiens histone deacetylase 3 (HDAC3), mRNA
NM 004964	Homo sapiens histone deacetylase 1 (HDAC1), mRNA
NM_001492	Homo sapiens growth differentiation factor 1 (GDF1), mRNA
NM_018486	Homo sapiens histone deacetylase 8 (HDAC8), mRNA
NM_005089	Homo sapiens U2 small nuclear ribonucleoprotein auxiliary factor, small subunit 2 (U2AF1RS2), mRNA
NM_004285	Homo sapiens hexose-6-phosphate dehydrogenase (glucose 1-dehydrogenase) (H6PD), mRNA
NM 007210	Homo sapiens UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-
	acetylgalactosaminyltransferase 6 (GalNAc-T6) (GALNT6), mRNA
NM 003774	Homo sapiens UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-
_	acetylgalactosaminyltransferase 4 (GalNAc-T4) (GALNT4), mRNA
NM_020474	Homo sapiens UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-
	acetylgalactosaminyltransferase 1 (GalNAc-T1) (GALNT1), mRNA
NM_015507	Homo sapiens EGF-like-domain, multiple 6 (EGFL6), mRNA
NM_004942	Homo sapiens defensin, beta 2 (DEFB2), mRNA
NM_005218	Homo sapiens defensin, beta 1 (DEFB1), mRNA
NM_002474	Homo sapiens myosin, heavy polypeptide 11, smooth muscle (MYH11),
	transcript variant SM1, mRNA
NM_022870	Homo sapiens myosin, heavy polypeptide 11, smooth muscle (MYH11),
	transcript variant SM3, mRNA
NM_022844	Homo sapiens myosin, heavy polypeptide 11, smooth muscle (MYH11),
	transcript variant SM2, mRNA
NM_001755	Homo sapiens core-binding factor, beta subunit (CBFB), transcript variant 2, mRNA
NM_016458	Homo sapiens hypothetical protein (LOC51236), mRNA
NM 020836	Homo sapiens KIAA1446 protein (KIAA1446), mRNA
NM 015407	Homo sapiens DKFZP564O243 protein (DKFZP564O243), mRNA
	- 17-10 DAY 21 3040243 protein (DAT 21 3040243), MRINA

	7771 10505 (CATA 40505) mPNA
NM_015062	Homo sapiens KIAA0595 protein (KIAA0595), mRNA
NM_019100	Homo sapiens DNA methyltransferase 1-associated protein 1 (DMAP1), mRNA
NM_015442	Homo sapiens hypothetical protein FLJ12890 (FLJ12890), mRNA  Homo sapiens hypothetical protein FLJ12890 (FLJ12890), mRNA
NM_023948	Homo sapiens hypothetical protein AF053356_CDS3 (AF053356_CDS3),
	mRNA
NM_022036	Homo sapiens G protein-coupled receptor, family C, group 5, member C
_	(GPRC5C), transcript variant 1, mRNA
NM 018653	Homo sapiens G protein-coupled receptor, family C, group 5, member C
-	(GPRC5C) transcript variant 2, mRNA
NM 000707	Homo sapiens arginine vasopressin receptor 1B (AVPR1B), mRNA
NM 000706	Homo saniens arginine vasopressin receptor IA (AVPRIA), mRNA
NM 021923	Homo senions fibroblast growth factor receptor-like 1 (FGFRL1), mRNA
NM 002011	Homo sapiens fibroblast growth factor receptor 4 (FGFR4), transcript variant 1,
11412_00_0	DNI 4
NM_022963	Homo sapiens fibroblast growth factor receptor 4 (FGFR4), transcript variant 2,
11117_0222	DNI A
NM 022965	Homo sapiens fibroblast growth factor receptor 3 (achondroplasia, thanatophoric
14141_022505	devention) (EGEP3) transcript variant 2, mRNA
NM 000142	Homo sapiens fibroblast growth factor receptor 3 (achondroplasia, thanatophoric
14141_000142	dwarfism) (FGFR3) transcript variant 1, mRNA
NM 022336	Homo sapiens ectodysplasin 1, anhidrotic receptor (EDAR), mRNA
NM_018654	Homo sapiens G protein-coupled receptor, family C, group 5, member D
MM_016034	(CDPC5D) mRNA
NM_002534	Homo sapiens 2',5'-oligoadenylate synthetase 1 (40-46 kD) (OAS1), transcript
NM_002334	variant E16 mDNA
ND 6 016016	Homo sapiens 2',5'-oligoadenylate synthetase 1 (40-46 kD) (OAS1), transcript
NM_016816	variant E18, mRNA
ND 6 014601	Homo sapiens ubiquitin carrier protein (E2-EPF), mRNA
NM_014501	Homo sapiens lymphotoxin alpha (TNF superfamily, member 1) (LTA), mRNA
NM_000595	Homo sapiens E1B-55kDa-associated protein 5 (E1B-AP5), mRNA
NM_007040	Homo sapiens calsequestrin 2 (cardiac muscle) (CASQ2), mRNA
NM_001232	Homo sapiens calsequestrin 2 (cardiac massle) (CASQ1), nuclear
NM_001231	gene encoding mitochondrial protein, mRNA
	Homo sapiens methyl-CpG binding domain protein 4 (MBD4), mRNA
NM_003925	Homo sapiens methyl-CpG binding domain protein 4 (AMDS 1), and the sapiens methyl-CpG binding domain protein 4 (AMDS 1), and the sapiens methyl-CpG binding domain protein 4 (AMDS 1), and the sapiens methyl-CpG binding domain protein 4 (AMDS 1), and the sapiens methyl-CpG binding domain protein 4 (AMDS 1), and the sapiens methyl-CpG binding domain protein 4 (AMDS 1), and the sapiens methyl-CpG binding domain protein 4 (AMDS 1), and the sapiens methyl-CpG binding domain protein 4 (AMDS 1), and the sapiens methyl-CpG binding domain protein 4 (AMDS 1), and the sapiens methyl-CpG binding domain protein 4 (AMDS 1), and the sapiens methyl-CpG binding domain protein 4 (AMDS 1), and the sapiens methyl-CpG binding domain protein 4 (AMDS 1), and the sapiens methyl-CpG binding domain protein 4 (AMDS 1), and the sapiens methyl-CpG binding domain protein 4 (AMDS 1), and the sapiens methyl-CpG binding domain protein 4 (AMDS 1), and the sapiens methyl-CpG binding domain protein 4 (AMDS 1), and the sapiens methyl-CpG binding domain bindin binding domain binding domain binding domain binding domain bin
NM_002059	Homo sapiens growth hormone 2 (GH2), transcript variant 1, mRNA
NM_022558	Homo sapiens growth hormone 2 (GH2), transcript variant 3, mRNA
NM_022557	Homo sapiens growth hormone 2 (GH2), transcript variant 2, mRNA  Homo sapiens growth hormone 2 (GH2), transcript variant 4, mRNA
NM_022556	Homo sapiens growth hormone 2 (GH2), transcript variant 4, mRNA
NM_022562	Homo sapiens growth hormone 1 (GH1), transcript variant 5, mRNA
NM_022561	Homo sapiens growth hormone 1 (GH1), transcript variant 4, mRNA
NM_022560	Homo sapiens growth hormone 1 (GH1), transcript variant 3, mRNA
NM 022559	Homo sapiens growth hormone 1 (GH1), transcript variant 2, mRNA
NM 000515	Homo sapiens growth hormone 1 (GH1), transcript variant 1, mRNA
NM 021801	Homo saniens matrix metalloproteinase 26 (MMP26), mRNA
NM 022718	Homo sapiens matrix metalloproteinase 25 (MMP25), transcript variant 2,
	mDNA
NM_022468	Homo sapiens matrix metalloproteinase 25 (MMP25), transcript variant 1,
11.1022.100	mPNA
NM 006690	Homo sapiens matrix metalloproteinase 24 (membrane-inserted) (MMP24),
1111_000000	mPNA
NM 004771	Homo saniens matrix metalloproteinase 20 (enamelysin) (MMP20), mRNA
	Homo sapiens matrix metalloproteinase 7 (matrilysin, uterine) (MMP7), mRNA
NM_002423	Homo suprens magny metanoprotestate , (

77.6.000.400	Type in a material constraint on 2 (otherwolvein 1 magazistings)
NM_002422	Homo sapiens matrix metalloproteinase 3 (stromelysin 1, progelatinase) (MMP3), mRNA
NM_005941	Homo sapiens matrix metalloproteinase 16 (membrane-inserted) (MMP16), transcript variant 1, mRNA
NM_022564	Homo sapiens matrix metalloproteinase 16 (membrane-inserted) (MMP16), transcript variant 2, mRNA
NM_002421	Homo sapiens matrix metalloproteinase 1 (interstitial collagenase) (MMP1), mRNA
NM_004995	Homo sapiens matrix metalloproteinase 14 (membrane-inserted) (MMP14), mRNA
NM 002427	Homo sapiens matrix metalloproteinase 13 (collagenase 3) (MMP13), mRNA
NM 005940	Homo sapiens matrix metalloproteinase 11 (stromelysin 3) (MMP11), mRNA
NM_022792	Homo sapiens matrix metalloproteinase 19 (MMP19), transcript variant rasi-9, mRNA
NM_022791	Homo sapiens matrix metalloproteinase 19 (MMP19), transcript variant rasi-6, mRNA
NM_022790	Homo sapiens matrix metalloproteinase 19 (MMP19), transcript variant rasi-3, mRNA
NM_002429	Homo sapiens matrix metalloproteinase 19 (MMP19), transcript variant rasi-1, mRNA
NM_004530	Homo sapiens matrix metalloproteinase 2 (gelatinase A, 72kD gelatinase, 72kD type IV collagenase) (MMP2), mRNA
NM_004994	Homo sapiens matrix metalloproteinase 9 (gelatinase B, 92kD gelatinase, 92kD type IV collagenase) (MMP9), mRNA
NM_004142	Homo sapiens matrix metalloproteinase-like 1 (MMPL1), mRNA
NM_002424	Homo sapiens matrix metalloproteinase 8 (neutrophil collagenase) (MMP8), mRNA
NM_002428	Homo sapiens matrix metalloproteinase 15 (membrane-inserted) (MMP15), mRNA
NM_002426	Homo sapiens matrix metalloproteinase 12 (macrophage elastase) (MMP12), mRNA
NM 002425	Homo sapiens matrix metalloproteinase 10 (stromelysin 2) (MMP10), mRNA
NM_022804	Homo sapiens SNRPN upstream reading frame (SNURF), transcript variant 2, mRNA
NM_005678	Homo sapiens SNRPN upstream reading frame (SNURF), transcript variant 1, mRNA
NM_003097	Homo sapiens small nuclear ribonucleoprotein polypeptide N (SNRPN), transcript variant 1, mRNA
NM_022808	Homo sapiens small nuclear ribonucleoprotein polypeptide N (SNRPN), transcript variant 5, mRNA
NM_022807	Homo sapiens small nuclear ribonucleoprotein polypeptide N (SNRPN), transcript variant 4, mRNA
NM_022806	Homo sapiens small nuclear ribonucleoprotein polypeptide N (SNRPN), transcript variant 3, mRNA
NM_022805	Homo sapiens small nuclear ribonucleoprotein polypeptide N (SNRPN), transcript variant 2, mRNA
NM_022717	Homo sapiens U1-snRNP binding protein homolog (70kD) (U1SNRNPBP), transcript variant 2, mRNA
NM_006759	Homo sapiens UDP-glucose pyrophosphorylase 2 (UGP2), mRNA
NM_001400	Homo sapiens endothelial differentiation, sphingolipid G-protein-coupled receptor, 1 (EDG1), mRNA
NM_005586	Homo sapiens MyoD family inhibitor (MDFI), mRNA

NM_022978	Homo sapiens small EDRK-rich factor 1B (centromeric) (SERF1B), mRNA
NM_023947	Homo sapiens hypothetical protein MGC3234 (MGC3234), mRNA
NM_023942	Homo sapiens hypothetical protein MGC3036 (MGC3036), mRNA
NM_023933	Homo sapiens hypothetical protein MGC2494 (MGC2494), mRNA
NM_005471	Homo sapiens glucosamine-6-phosphate isomerase (GNPI), mRNA
NM_023925	Homo sapiens hypothetical protein FLJ22569 (FLJ22569), mRNA
NM 004076	Homo sapiens crystallin, beta B3 (CRYBB3), mRNA
NM 015717	Homo sapiens Langerhans cell specific c-type lectin (LANGERIN), mRNA
NM_012329	Homo sapiens monocyte to macrophage differentiation-associated (MMD), mRNA
NM_007020	Homo sapiens U1-snRNP binding protein homolog (70kD) (U1SNRNPBP), transcript variant 1, mRNA
NM_006465	Homo sapiens dead ringer (Drosophila)-like 2 (bright and dead ringer) (DRIL2), mRNA
NM_000015	Homo sapiens N-acetyltransferase 2 (arylamine N-acetyltransferase) (NAT2), mRNA
NM_000496	Homo sapiens crystallin, beta B2 (CRYBB2), mRNA
NM_001886	Homo sapiens crystallin, beta A4 (CRYBA4), mRNA
NM_023080	Homo sapiens hypothetical protein FLJ20989 (FLJ20989), mRNA
NM 023039	Homo sapiens ankyrin repeat, family A (RFXANK-like), 2 (ANKRA2), mRNA
NM_021905	Homo sapiens gamma-aminobutyric acid (GABA) B receptor, 1 (GABBR1), transcript variant 4, mRNA
NM_020554	Homo sapiens T-cell leukemia/lymphoma 6 (TCL6), transcript variant TCL6d1, mRNA
NM_020553	Homo sapiens T-cell leukemia/lymphoma 6 (TCL6), transcript variant TCL6c1, mRNA
NM_020552	Homo sapiens T-cell leukemia/lymphoma 6 (TCL6), transcript variant TCL6b1, mRNA
NM_020550	Homo sapiens T-cell leukemia/lymphoma 6 (TCL6), transcript variant TCL6a3, mRNA
NM_012468	Homo sapiens T-cell leukemia/lymphoma 6 (TCL6), transcript variant TCL6a1, mRNA
NM_014418	Homo sapiens T-cell leukemia/lymphoma 6 (TCL6), transcript variant TCL6a2, mRNA
NM 016730	Homo sapiens folate receptor 1 (adult) (FOLR1), transcript variant 3, mRNA
NM 016729	Homo sapiens folate receptor 1 (adult) (FOLR1), transcript variant 4, mRNA
NM 016725	Homo sapiens folate receptor 1 (adult) (FOLR1), transcript variant 1, mRNA
NM 016724	Homo sapiens folate receptor 1 (adult) (FOLR1), transcript variant 7, mRNA
NM 016025	Homo sapiens CGI-81 protein (DREV1), mRNA
NM_004406	Homo sapiens deleted in malignant brain tumors 1 (DMBT1), transcript variant 1, mRNA
NM 000197	Homo sapiens hydroxysteroid (17-beta) dehydrogenase 3 (HSD17B3), mRNA
NM_001220	Homo sapiens calcium/calmodulin-dependent protein kinase (CaM kinase) II
1414_001220	beta (CAMK2B), mRNA
NM 019071	Homo sapiens inhibitor of growth family, member 3 (ING3), mRNA
NM 016731	Homo sapiens folate receptor 1 (adult) (FOLR1), transcript variant 8, mRNA
NM 023018	Homo sapiens hypothetical protein FLJ13052 (FLJ13052), mRNA
NM 023016	Homo sapiens hypothetical protein FLJ21870 (FLJ21870), mRNA
NM 022911	Homo sapiens solute carrier family 26, member 6 (SLC26A6), mRNA
NM 021071	Homo sapiens ADP-ribosyltransferase 4 (ART4), mRNA
NM 022113	Homo sapiens kinesin family member 13A (KIF13A), mRNA
NM 012449	Homo sapiens six transmembrane epithelial antigen of the prostate (STEAP),
14141 012443	Alone especia dia transmissione epitteria mangana and procession (2000)

	mRNA
NM 016513	Homo sapiens MAK-related kinase (KIAA0936), mRNA
NM 014920	Homo sapiens MAK-related kinase (KIAA0936), mRNA
NM 014688	Homo sapiens related to the N terminus of tre (RNTRE), mRNA
NM 006640	Homo sapiens MLL septin-like fusion (MSF), mRNA
NM 006070	Homo sapiens TRK-fused gene (TFG), mRNA
NM 004809	Homo sapiens stomatin-like 1 (STOML1), mRNA
NM 000297	Homo sapiens polycystic kidney disease 2 (autosomal dominant) (PKD2),
11111	mRNA
NM 016307	Homo sapiens paired related homeobox protein (PRX2), mRNA
NM 003924	Homo sapiens paired mesoderm homeobox 2b (PMX2B), mRNA
NM_006902	Homo sapiens paired mesoderm homeo box 1 (PMX1), transcript variant pmx-1a, mRNA
NM_022716	Homo sapiens paired mesoderm homeo box 1 (PMX1), transcript variant pmx-1b, mRNA
NM_000916	Homo sapiens oxytocin receptor (OXTR), mRNA
NM_000915	Homo sapiens oxytocin, prepro- (neurophysin I) (OXT), mRNA
NM_006188	Homo sapiens oncomodulin (OCM), mRNA
NM_022664	Homo sapiens extracellular matrix protein 1 (ECM1), transcript variant 2, mRNA
NM_004092	Homo sapiens enoyl Coenzyme A hydratase, short chain, 1, mitochondrial (ECHS1), nuclear gene encoding mitochondrial protein, mRNA
NM_022652	Homo sapiens dual specificity phosphatase 6 (DUSP6), transcript variant 2, mRNA
NM_004419	Homo sapiens dual specificity phosphatase 5 (DUSP5), mRNA
NM_004425	Homo sapiens extracellular matrix protein 1 (ECM1), transcript variant 1, mRNA
NM_004418	Homo sapiens dual specificity phosphatase 2 (DUSP2), mRNA
NM_004961	Homo sapiens gamma-aminobutyric acid (GABA) A receptor, epsilon (GABRE), transcript variant 1, mRNA
NM_021990	Homo sapiens gamma-aminobutyric acid (GABA) A receptor, epsilon (GABRE), transcript variant 4, mRNA
NM_021987	Homo sapiens gamma-aminobutyric acid (GABA) A receptor, epsilon (GABRE), transcript variant 3, mRNA
NM_021984	Homo sapiens gamma-aminobutyric acid (GABA) A receptor, epsilon (GABRE), transcript variant 2, mRNA
NM_004090	Homo sapiens dual specificity phosphatase 3 (vaccinia virus phosphatase VH1-related) (DUSP3), mRNA
NM_001398	Homo sapiens enoyl Coenzyme A hydratase 1, peroxisomal (ECH1), mRNA
NM_001946	Homo sapiens dual specificity phosphatase 6 (DUSP6), transcript variant 1, mRNA
NM_001952	Homo sapiens E2F transcription factor 6 (E2F6), mRNA
NM 001950	Homo sapiens E2F transcription factor 4, p107/p130-binding (E2F4), mRNA
NM_001949	Homo sapiens E2F transcription factor 3 (E2F3) mRNA, complete cds
NM_005225	Homo sapiens E2F transcription factor 1 (E2F1), mRNA
NM_022977	Homo sapiens fatty-acid-Coenzyme A ligase, long-chain 4 (FACL4), transcript variant 2, mRNA
NM_004457	Homo sapiens fatty-acid-Coenzyme A ligase, long-chain 3 (FACL3), mRNA
NM_021122	Homo sapiens fatty-acid-Coenzyme A ligase, long-chain 2 (FACL2), mRNA
NM_002473	Homo sapiens myosin, heavy polypeptide 9, non-muscle (MYH9), mRNA
NM_001926	Homo sapiens defensin, alpha 6, Paneth cell-specific (DEFA6), mRNA
NM_005217	Homo sapiens defensin, alpha 3, neutrophil-specific (DEFA3), mRNA

NM_021912	Homo sapiens gamma-aminobutyric acid (GABA) A receptor, beta 3
	(GABRB3), transcript variant 2, mRNA
NM_021911	Homo sapiens gamma-aminobutyric acid (GABA) A receptor, beta 2 (GABRB2), transcript variant 1, mRNA
ND 4 000014	Homo sapiens gamma-aminobutyric acid (GABA) A receptor, beta 3
NM_000814	(GABRB3), transcript variant 1, mRNA
NM_000812	Homo sapiens gamma-aminobutyric acid (GABA) A receptor, beta 1
	(GABRB1), mRNA
NM_022650	Homo sapiens RAS p21 protein activator (GTPase activating protein) 1
_	(RASA1), transcript variant 2, mRNA
NM 003259	Homo sapiens intercellular adhesion molecule 5, telencephalin (ICAM5), mRNA
NM 022377	Homo sapiens intercellular adhesion molecule 4, Landsteiner-Wiener blood
1111_022577	group (ICAM4), transcript variant 2, mRNA
NM 001544	Homo sapiens intercellular adhesion molecule 4, Landsteiner-Wiener blood
14141_0015-1-1	group (ICAM4), transcript variant 1, mRNA
NM 002162	Homo sapiens intercellular adhesion molecule 3 (ICAM3), mRNA
NM 000873	Homo sapiens intercellular adhesion molecule 2 (ICAM2), mRNA
NM 022308	Homo sapiens islet cell autoantigen 1 (69kD) (ICA1), transcript variant 3,
NWI_022308	mRNA
NM_022307	Homo sapiens islet cell autoantigen 1 (69kD) (ICA1), transcript variant 1,
	mRNA
NM_022581	Homo sapiens chorionic somatomammotropin hormone-like 1 (CSHL1),
	transcript variant 5, mRNA
NM_022580	Homo sapiens chorionic somatomammotropin hormone-like 1 (CSHL1),
1444_022500	transcript variant 4, mRNA
NM_022579	Homo sapiens chorionic somatomammotropin hormone-like 1 (CSHL1),
14141_022379	transcript variant 3, mRNA
NM_022578	Homo sapiens chorionic somatomammotropin hormone-like 1 (CSHL1),
1111_022576	transcript variant 2, mRNA
NM_001318	Homo sapiens chorionic somatomammotropin hormone-like 1 (CSHL1),
MM_001318	transcript variant 1, mRNA
NM 022646	Homo sapiens chorionic somatomammotropin hormone 2 (CSH2), transcript
NWI_022046	variant 4, mRNA
NM 022645	Homo sapiens chorionic somatomammotropin hormone 2 (CSH2), transcript
NIVI_022643	variant 3, mRNA
ND 6 000644	
NM_022644	Homo sapiens chorionic somatomammotropin hormone 2 (CSH2), transcript
12 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2	variant 2, mRNA
NM_020991	Homo sapiens chorionic somatomammotropin hormone 2 (CSH2), transcript variant 1, mRNA
NM 022642	Homo sapiens chorionic somatomammotropin hormone 1 (placental lactogen)
NW1_022042	(CSH1), transcript variant 4, mRNA
NM 022641	Homo sapiens chorionic somatomammotropin hormone 1 (placental lactogen)
NIVI_022041	(CSH1), transcript variant 3, mRNA
ND 6 022640	Homo sapiens chorionic somatomammotropin hormone 1 (placental lactogen)
NM_022640	
377.6.001015	(CSH1), transcript variant 2, mRNA
NM_001317	Homo sapiens chorionic somatomammotropin hormone 1 (placental lactogen)
	(CSH1), transcript variant 1, mRNA
NM_002371	Homo sapiens mal, T-cell differentiation protein (MAL), transcript variant a,
	mRNA
NM_022440	Homo sapiens mal, T-cell differentiation protein (MAL), transcript variant d,
	mRNA
NM_022439	Homo sapiens mal, T-cell differentiation protein (MAL), transcript variant c,

	L DAYA
NR 6 000 100	mRNA
NM_022438	Homo sapiens mal, T-cell differentiation protein (MAL), transcript variant b,
37.6 001500	mRNA
NM_001790	Homo sapiens cell division cycle 25C (CDC25C), transcript variant 1, mRNA
NM_022809	Homo sapiens cell division cycle 25C (CDC25C), transcript variant 2, mRNA
NM_021141	Homo sapiens X-ray repair complementing defective repair in Chinese hamster
27.5 000550	cells 5 (double-strand-break rejoining; Ku autoantigen, 80kD) (XRCC5), mRNA
NM_022550	Homo sapiens X-ray repair complementing defective repair in Chinese hamster
NDM 022406	cells 4 (XRCC4), transcript variant 3, mRNA  Homo sapiens X-ray repair complementing defective repair in Chinese hamster
NM_022406	cells 4 (XRCC4), transcript variant 2, mRNA
NM 005432	Homo sapiens X-ray repair complementing defective repair in Chinese hamster
NWI_003432	cells 3 (XRCC3), mRNA
NM_003401	Homo sapiens X-ray repair complementing defective repair in Chinese hamster
11111_005401	cells 4 (XRCC4), transcript variant 1, mRNA
NM 022405	Homo sapiens X transporter protein 3 (XT3), transcript variant 2, mRNA
NM 016192	Homo sapiens transporter protein with EGF-like and two follistatin-like
1411_010172	domains 2 (TMEFF2), mRNA
NM 006786	Homo sapiens urotensin 2 (UTS2), transcript variant 2, mRNA
NM 021995	Homo sapiens urotensin 2 (UTS2), transcript variant 1, mRNA
NM 003353	Homo sapiens urocortin (UCN), mRNA
NM 021991	Homo sapiens junction plakoglobin (JUP), transcript variant 2, mRNA
NM 021737	Homo sapiens chloride channel 6 (CLCN6), transcript variant ClC-6d, mRNA.
NM 021736	Homo sapiens chloride channel 6 (CLCN6), transcript variant ClC-6c, mRNA
NM 021735	Homo sapiens chloride channel 6 (CLCN6), transcript variant ClC-6b, mRNA
NM 006536	Homo sapiens chloride channel, calcium activated, family member 2 (CLCA2),
	mRNA
NM 004000	Homo sapiens chitinase 3-like 2 (CHI3L2), mRNA
NM 002641	Homo sapiens phosphatidylinositol glycan, class A (paroxysmal nocturnal
_	hemoglobinuria) (PIGA), transcript variant 1, mRNA
NM_020473	Homo sapiens phosphatidylinositol glycan, class A (paroxysmal nocturnal
	hemoglobinuria) (PIGA), transcript variant 3, mRNA
NM_020472	Homo sapiens phosphatidylinositol glycan, class A (paroxysmal nocturnal
	hemoglobinuria) (PIGA), transcript variant 2, mRNA
NM_001699	Homo sapiens AXL receptor tyrosine kinase (AXL), transcript variant 2, mRNA
NM_021913	Homo sapiens AXL receptor tyrosine kinase (AXL), transcript variant 1, mRNA
NM_016188	Homo sapiens actin-like 6 (ACTL6), mRNA
NM_000509	Homo sapiens fibrinogen, gamma polypeptide (FGG), transcript variant gamma-
77.7	A, mRNA
NM_021870	Homo sapiens fibrinogen, gamma polypeptide (FGG), transcript variant gamma-
77.5.6055115	B, mRNA
NM_005141	Homo sapiens fibrinogen, B beta polypeptide (FGB), mRNA
NM_021871	Homo sapiens fibrinogen, A alpha polypeptide (FGA), transcript variant alpha,
NIM DODGOO	mRNA  Home senions fibringers A alpha palamentide (EGA) transcript verient alpha E
NM_000508	Homo sapiens fibrinogen, A alpha polypeptide (FGA), transcript variant alpha-E, mRNA
NM 000920	Homo sapiens pyruvate carboxylase (PC), nuclear gene encoding mitochondrial
11112_000,20	protein, transcript variant A, mRNA
NM 022172	Homo sapiens pyruvate carboxylase (PC), nuclear gene encoding mitochondrial
	protein, transcript variant 2, mRNA
NM 004358	Homo sapiens cell division cycle 25B (CDC25B), transcript variant 1, mRNA
NM 021874	Homo sapiens cell division cycle 25B (CDC25B), transcript variant 4, mRNA

	1 25D (CDC25D) +int 2 DNA
NM_021873	Homo sapiens cell division cycle 25B (CDC25B), transcript variant 3, mRNA
NM_021872	Homo sapiens cell division cycle 25B (CDC25B), transcript variant 2, mRNA
NM_020990	Homo sapiens creatine kinase, mitochondrial 1 (ubiquitous) (CKMT1), nuclear
	gene encoding mitochondrial protein, mRNA
NM_021962	Homo sapiens active BCR-related gene (ABR), transcript variant 1, mRNA
NM_001092	Homo sapiens active BCR-related gene (ABR), transcript variant 2, mRNA
NM_021794	Homo sapiens a disintegrin and metalloproteinase domain 30 (ADAM30), transcript variant 1, mRNA
NM_001464	Homo sapiens a disintegrin and metalloproteinase domain 2 (fertilin beta)
NM_021780	Homo sapiens a disintegrin and metalloproteinase domain 29 (ADAM29), transcript variant 2, mRNA
NM_021779	Homo sapiens a disintegrin and metalloproteinase domain 29 (ADAM29), transcript variant 3, mRNA
NM_014269	Homo sapiens a disintegrin and metalloproteinase domain 29 (ADAM29), transcript variant 1, mRNA
NM_021723	Homo sapiens a disintegrin and metalloproteinase domain 22 (ADAM22), mRNA
NM_021722	Homo sapiens a disintegrin and metalloproteinase domain 22 (ADAM22), mRNA
NM_021721	Homo sapiens a disintegrin and metalloproteinase domain 22 (ADAM22), mRNA
NM_016351	Homo sapiens a disintegrin and metalloproteinase domain 22 (ADAM22), mRNA
NM_021832	Homo sapiens a disintegrin and metalloproteinase domain 17 (tumor necrosis factor, alpha, converting enzyme) (ADAM17), transcript variant 2, mRNA
NM_003183	Homo sapiens a disintegrin and metalloproteinase domain 17 (tumor necrosis factor, alpha, converting enzyme) (ADAM17), transcript variant 1, mRNA
NM_003815	Homo sapiens a disintegrin and metalloproteinase domain 15 (metargidin) (ADAM15), mRNA
NM_021641	Homo sapiens a disintegrin and metalloproteinase domain 12 (meltrin alpha) (ADAM12), transcript variant 2, mRNA
NM_021612	Homo sapiens a disintegrin and metalloproteinase domain 11 (ADAM11), transcript variant 2, mRNA
NM_006437	Homo sapiens ADP-ribosyltransferase (NAD+; poly (ADP-ribose) polymerase)-
NM_001618	Homo sapiens ADP-ribosyltransferase (NAD+; poly (ADP-ribose) polymerase) (ADPRT), mRNA
NM 021738	Homo sapiens supervillin (SVIL), transcript variant 2, mRNA
NM 003174	Homo sapiens supervillin (SVIL), transcript variant 1, mRNA
NM_002505	Homo sapiens nuclear transcription factor Y, alpha (NFYA), transcript variant 1, mRNA
NM_021705	Homo sapiens nuclear transcription factor Y, alpha (NFYA), transcript variant 2, mRNA
NM_000832	Homo sapiens glutamate receptor, ionotropic, N-methyl D-aspartate 1 (GRIN1), transcript variant NR1-1, mRNA
NM_000673	Homo sapiens alcohol dehydrogenase 7 (class IV), mu or sigma polypeptide (ADH7), mRNA
NM_000671	Homo sapiens alcohol dehydrogenase 5 (class III), chi polypeptide (ADH5), mRNA
NM_000670	Homo sapiens alcohol dehydrogenase 4 (class II), pi polypeptide (ADH4), mRNA

Times and and additional managestic (CLDC) DNIA
Homo sapiens colipase, pancreatic (CLPS), mRNA
Homo sapiens ELK4, ETS-domain protein (SRF accessory protein 1) (ELK4),
transcript variant b, mRNA
Homo sapiens CD27-binding (Siva) protein (SIVA), transcript variant 2, mRNA
Homo sapiens CD27-binding (Siva) protein (SIVA), transcript variant 1, mRNA
Homo sapiens angiotensin I converting enzyme (peptidyl-dipeptidase A) 2 (ACE2), mRNA
Homo sapiens X transporter protein 3 (XT3), transcript variant 1, mRNA
Homo sapiens zinc finger protein 14 (KOX 6) (ZNF14), mRNA
Homo sapiens Rhesus blood group, CcEe antigens (RHCE), mRNA
Homo sapiens interleukin 1 receptor-like 1 (IL1RL1), mRNA
Homo sapiens FXYD domain-containing ion transport regulator 2 (FXYD2),
transcript variant a, mRNA
Homo sapiens FXYD domain-containing ion transport regulator 2 (FXYD2),
transcript variant b, mRNA
Homo sapiens nucleoporin 98kD (NUP98), mRNA
Homo sapiens CD79B antigen (immunoglobulin-associated beta) (CD79B),
transcript variant 2, mRNA
Homo sapiens CD79B antigen (immunoglobulin-associated beta) (CD79B),
transcript variant 1, mRNA
Homo sapiens CD79A antigen (immunoglobulin-associated alpha) (CD79A),
transcript variant 2, mRNA
Homo sapiens a disintegrin-like and metalloprotease (reprolysin type) with
thrombospondin type 1 motif, 2 (ADAMTS2), transcript variant 2, mRNA
Homo sapiens a disintegrin-like and metalloprotease (reprolysin type) with
thrombospondin type 1 motif, 1 (ADAMTS1), mRNA
Homo sapiens adaptor-related protein complex 2, sigma 1 subunit (AP2S1),
transcript variant AP17, mRNA
Homo sapiens adaptor-related protein complex 2, sigma 1 subunit (AP2S1),
transcript variant AP17delta, mRNA
Homo sapiens breakpoint cluster region (BCR), transcript variant 2, mRNA
Homo sapiens breakpoint cluster region (BCR), transcript variant 1, mRNA
Homo sapiens glutamate receptor, ionotropic, N-methyl D-aspartate 1 (GRIN1), transcript variant NR1-3, mRNA
Homo sapiens glutamate receptor, ionotropic, N-methyl D-aspartate 1 (GRIN1),
transcript variant NR1-2, mRNA
Homo sapiens choline acetyltransferase (CHAT), transcript variant R, mRNA
Homo sapiens choline acetyltransferase (CHAT), transcript variant N1, mRNA
Homo sapiens choline acetyltransferase (CHAT), transcript variant M, mRNA
Homo sapiens actin, gamma 2, smooth muscle, enteric (ACTG2), mRNA
Homo sapiens choline acetyltransferase (CHAT), transcript variant N2, mRNA
Homo sapiens disrupted in schizophrenia 1 (DISC1), mRNA
Homo sapiens mucin 4, tracheobronchial (MUC4), mRNA
Homo sapiens hypothetical protein FLJ20357 (FLJ20357), mRNA
Homo sapiens mucin 4, tracheobronchial (MUC4), mRNA
Homo sapiens meningioma expressed antigen 5 (hyaluronidase) (MGEA5),
mRNA
Homo sapiens ATP-binding cassette, sub-family D (ALD), member 4 (ABCD4),
transcript variant 5, mRNA
Homo sapiens ATP-binding cassette, sub-family D (ALD), member 4 (ABCD4),
transcript variant 4, mRNA
Homo sapiens ATP-binding cassette, sub-family D (ALD), member 4 (ABCD4),

	L. C. L. C. L. PNIA
	transcript variant 3, mRNA
NM_020323	Homo sapiens ATP-binding cassette, sub-family D (ALD), member 4 (ABCD4),
	transcript variant 2, mRNA
NM_020298	Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 9
	(ABCC9), transcript variant SUR2A-delta-14, mRNA
NM_020297	Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 9
	(ABCC9), transcript variant SUR2B, mRNA
NM_021270	Homo sapiens leukocyte-associated Ig-like receptor 2 (LAIR2), transcript variant
_	2. mRNA
NM_002288	Homo sapiens leukocyte-associated Ig-like receptor 2 (LAIR2), transcript variant
	1, mRNA
NM 020983	Homo sapiens adenylate cyclase 6 (ADCY6), transcript variant 2, mRNA
NM 015270	Homo sapiens adenylate cyclase 6 (ADCY6), transcript variant 1, mRNA
NM 020987	Homo sapiens ankyrin 3, node of Ranvier (ankyrin G) (ANK3), transcript variant
14W1_020987	1, mRNA
NM 020977	Homo sapiens ankyrin 2, neuronal (ANK2), transcript variant 2, mRNA
	Homo sapiens ankyrin 2, neuronal (ANK2), transcript variant 1, mRNA
NM_001148	TY and the state of the state o
NM_020481	Homo sapiens ankyrin 1, erythrocytic (ANK1), transcript variant 8, mRNA
NM_020480	Homo sapiens ankyrin 1, erythrocytic (ANK1), transcript variant 7, mRNA
NM_020479	Homo sapiens ankyrin 1, erythrocytic (ANK1), transcript variant 6, mRNA
NM_020478	Homo sapiens ankyrin 1, erythrocytic (ANK1), transcript variant 5, mRNA
NM_020477	Homo sapiens ankyrin 1, erythrocytic (ANK1), transcript variant 2, mRNA
NM 000037	Homo sapiens ankyrin 1, erythrocytic (ANK1), transcript variant 3, mRNA
NM 020476	Homo sapiens ankyrin 1, erythrocytic (ANK1), transcript variant 1, mRNA
NM 020475	Homo sapiens ankyrin 1, erythrocytic (ANK1), transcript variant 4, mRNA
NM 021056	Homo sapiens tuberous sclerosis 2 (TSC2), transcript variant 3, mRNA
NM 021055	Homo sapiens tuberous sclerosis 2 (TSC2), transcript variant 2, mRNA
NM 000548	Homo sapiens tuberous sclerosis 2 (TSC2), transcript variant 1, mRNA
NM 004041	Homo sapiens arrestin, beta 1 (ARRB1), transcript variant 1, mRNA
NM 020251	Homo sapiens arrestin, beta 1 (ARRB1), transcript variant 2, mRNA
NM_000872	Homo sapiens 5-hydroxytryptamine (serotonin) receptor 7 (adenylate cyclase-
19191_000872	coupled) (HTR7), transcript variant a, mRNA
ND ( 010000	Homo sapiens 5-hydroxytryptamine (serotonin) receptor 7 (adenylate cyclase-
NM_019860	Homo sapiens 5-nydroxytryptamine (serotomin) receptor / (adenylate cyclase-
37.5.010050	coupled) (HTR7), transcript variant b, mRNA
NM_019859	Homo sapiens 5-hydroxytryptamine (serotonin) receptor 7 (adenylate cyclase-
	coupled) (HTR7), transcript variant d, mRNA
NM_004228	Homo sapiens pleckstrin homology, Sec7 and coiled/coil domains 2 (cytohesin-
	2) (PSCD2), transcript variant 2, mRNA
NM_017457	Homo sapiens pleckstrin homology, Sec7 and coiled/coil domains 2 (cytohesin-
	2) (PSCD2), transcript variant 1, mRNA
NM_004302	Homo sapiens activin A receptor, type IB (ACVR1B), transcript variant 1,
	mRNA
NM_020328	Homo sapiens activin A receptor, type IB (ACVR1B), transcript variant 3,
_	mRNA
NM 020327	Homo sapiens activin A receptor, type IB (ACVR1B), transcript variant 2,
	mRNA
NM 012082	Homo sapiens Friend of GATA2 (FOG2), mRNA
NM 000578	Homo sapiens solute carrier family 11 (proton-coupled divalent metal ion
1 000578	transporters), member 1 (SLC11A1), mRNA
NM 021094	Homo sapiens solute carrier family 21 (organic anion transporter), member 3
14141_021094	(SLC21A3), mRNA
NIM 002720	Homo sapiens aldo-keto reductase family 1, member C3 (3-alpha hydroxysteroid
NM_003739	nomo sapiens aido-keio reduciase family 1, member 65 (5-alpha nydroxysteroid

	dehydrogenase, type II) (AKR1C3), mRNA
NM_000735	Homo sapiens glycoprotein hormones, alpha polypeptide (CGA), mRNA
NM_014272	Homo sapiens a disintegrin-like and metalloprotease (reprolysin type) with
	thrombospondin type 1 motif, 7 (ADAMTS7), mRNA
NM_019863	Homo sapiens coagulation factor VIII, procoagulant component (hemophilia A)
	(F8), transcript variant 2, mRNA
NM_000132	Homo sapiens coagulation factor VIII, procoagulant component (hemophilia A) (F8), transcript variant 1, mRNA
NM_019616	Homo sapiens coagulation factor VII (serum prothrombin conversion
	accelerator) (F7), transcript variant 2, mRNA
NM_000131	Homo sapiens coagulation factor VII (serum prothrombin conversion
	accelerator) (F7), transcript variant 1, mRNA
NM_007219	Homo sapiens ring finger protein 24 (RNF24), mRNA
NM_021010	Homo sapiens defensin, alpha 5, Paneth cell-specific (DEFA5), mRNA
NM_016250	Homo sapiens N-myc downstream-regulated gene 2 (NDRG2), mRNA
NM 020525	Homo sapiens interleukin 22 (IL22), mRNA
NM 006774	Homo sapiens indolethylamine N-methyltransferase (INMT), mRNA
NM 014310	Homo sapiens similar to mouse Ras, dexamethasone-induced 1 (RASD1),
	mRNA
NM_020322	Homo sapiens amiloride-sensitive cation channel 3, testis (ACCN3), transcript variant 3, mRNA
NM_020321	Homo sapiens amiloride-sensitive cation channel 3, testis (ACCN3), transcript
	variant 2, mRNA
NM_020334	Homo sapiens a disintegrin and metalloproteinase domain 30 (ADAM30),
. <del>-</del>	transcript variant 2, mRNA
NM_019559	Homo sapiens coagulation factor XI (plasma thromboplastin antecedent) (F11),
_	transcript variant 2, mRNA
NM_000128	Homo sapiens coagulation factor XI (plasma thromboplastin antecedent) (F11),
	transcript variant 1, mRNA
NM_000443	Homo sapiens ATP-binding cassette, sub-family B (MDR/TAP), member 4 (ABCB4), transcript variant A, mRNA
NM_018850	Homo sapiens ATP-binding cassette, sub-family B (MDR/TAP), member 4
	(ABCB4), transcript variant C, mRNA
NM_018849	Homo sapiens ATP-binding cassette, sub-family B (MDR/TAP), member 4
	(ABCB4), transcript variant B, mRNA
NM_020038	Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 3
	(ABCC3), transcript variant MRP3B, mRNA
NM_020037	Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 3
	(ABCC3), transcript variant MRP3A, mRNA
NM_003786	Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 3
_	(ABCC3), transcript variant MRP3, mRNA
NM 019624	Homo sapiens ATP-binding cassette, sub-family B (MDR/TAP), member 9
 I	(ABCB9), transcript variant 2, mRNA
NM 019625	Homo sapiens ATP-binding cassette, sub-family B (MDR/TAP), member 9
_	(ABCB9), transcript variant 1, mRNA
NM 004996	Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 1
_	(ABCC1), transcript variant 1, mRNA
NM 019902	Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 1
<del>-</del> * * * * * * * * * * * * * * * * * * *	(ABCC1), transcript variant 7, mRNA
NM 019901	Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 1
_ "	
	(ABCC1), transcript variant 6, mRNA

	(ADCCI) / which might 5 mpDNA
	(ABCC1), transcript variant 5, mRNA
NM_019899	Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 1
	(ABCC1), transcript variant 4, mRNA  Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 1
NM_019898	Homo sapiens ATP-binding cassette, sub-family C (CI Tioving), moment
37.6.010060	(ABCC1), transcript variant 3, mRNA  Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 1
NM_019862	Homo sapiens ATP-binding casselle, sub-laminy C (CI Titolving), member 1
,	(ABCC1), transcript variant 2, mRNA
NM_019903	Homo sapiens adducin 3 (gamma) (ADD3), transcript variant 2, mRNA
NM_001640	Homo sapiens N-acylaminoacyl-peptide hydrolase (APEH), mRNA
NM_019858	Homo sapiens protein A (A), transcript variant A-2, mRNA
NM_000407	Homo sapiens glycoprotein Ib (platelet), beta polypeptide (GP1BB), mRNA
NM_015675	Homo sapiens growth arrest and DNA-damage-inducible, beta (GADD45B), mRNA
NM 016824	Homo sapiens adducin 3 (gamma) (ADD3), transcript variant 1, mRNA
NM_020039	Homo sapiens amiloride-sensitive cation channel 2, neuronal (ACCN2),
NT 6 005000	transcript variant 1, mRNA
NM_005388	Homo sapiens phosducin-like (PDCL), mRNA  Homo sapiens solute carrier family 2 (facilitated glucose transporter), member 6
NM_017585	(SLC2A6) mRNA
NM 020238	Homo sapiens inner centromere protein antigens (135kD, 155kD) (INCENP),
_	mRNA
NM 006908	Homo sapiens ras-related C3 botulinum toxin substrate 1 (rho family, small GTP
_	binding protein Rac1) (RAC1), transcript variant Rac1, mRNA
NM_018890	Homo sapiens ras-related C3 botulinum toxin substrate 1 (rho family, small GTP
	binding protein Rac1) (RAC1), transcript variant Rac1b, mRNA
NM 018891	Homo sapiens laminin, gamma 2 (nicein (100kD), kalinin (105kD), BM600
<u></u>	(100kD), Herlitz junctional epidermolysis bullosa)) (LAMC2), transcript variant 2, mRNA
NM_013430	Homo sapiens gamma-glutamyltransferase 1 (GGT1), transcript variant 3,
14141_013430	mRNA
NM_013421	Homo sapiens gamma-glutamyltransferase 1 (GGT1), transcript variant 2,
NWI_013421	mRNA
NM 004954	Homo sapiens ELKL motif kinase (EMK1), transcript variant 2, mRNA
NM 017490	Homo sapiens ELKL motif kinase (EMK1), transcript variant 1, mRNA
NM 004105	Homo sapiens EGF-containing fibulin-like extracellular matrix protein 1
14141_004103	(FFFMP1) transcript variant 1, mRNA
NM 002403	Homo sapiens microfibrillar-associated protein 2 (MFAP2), transcript variant 2,
	mRNA
NM 017459	Homo sapiens microfibrillar-associated protein 2 (MFAP2), transcript variant 1,
	mRNA
NM 005115	Homo sapiens major vault protein (MVP), transcript variant 2, mRNA
NM 017458	Homo sapiens major vault protein (MVP), transcript variant 1, mRNA
NM 018894	Homo sapiens EGF-containing fibulin-like extracellular matrix protein 1
	(EFEMP1), transcript variant 2, mRNA
NM 016519	Homo sapiens ameloblastin, enamel matrix protein (AMBN), mRNA
NM 017492	Homo sapiens ataxin 2 related protein (A2LP), transcript variant 2, mRNA
NM 007193	Homo sapiens annexin A10 (ANXA10), mRNA
NM 019102	Homo sapiens homeo box A5 (HOXA5), mRNA
NM 018971	Homo sapiens G protein-coupled receptor 27 (GPR27), mRNA
NM 003379	Homo sapiens villin 2 (ezrin) (VIL2), mRNA
	1
NM 016830	Homo sapiens vesicle-associated membrane protein 1 (synaptobrevin 1)

NM_014231	Homo sapiens vesicle-associated membrane protein 1 (synaptobrevin 1)
	(VAMP1), transcript variant VAMP-1A, mRNA
NM_017489	Homo sapiens telomeric repeat binding factor (NIMA-interacting) 1 (TERF1), transcript variant 1, mRNA
NT 6 002210	Homo sapiens telomeric repeat binding factor (NIMA-interacting) 1 (TERF1),
NM_003218	transcript variant 2, mRNA
NM_017455	Homo sapiens stromal cell derived factor receptor 1 (SDFR1), transcript variant
	alpha, mRNA
NM_007098	Homo sapiens clathrin, heavy polypeptide-like 1 (CLTCL1), transcript variant 2,
	mRNA
NM 017451	Homo sapiens BAI1-associated protein 2 (BAIAP2), transcript variant 2, mRNA
NM 017450	Homo sapiens BAI1-associated protein 2 (BAIAP2), transcript variant 1, mRNA
NM 001617	Homo sapiens adducin 2 (beta) (ADD2), transcript variant beta-1, mRNA
NM 017488	Homo sapiens adducin 2 (beta) (ADD2), transcript variant beta-4, mRNA
NM 017487	Homo sapiens adducin 2 (beta) (ADD2), transcript variant beta-6b, mRNA
NM 017486	Homo sapiens adducin 2 (beta) (ADD2), transcript variant beta-6a, mRNA
NM 017485	Homo sapiens adducin 2 (beta) (ADD2), transcript variant beta-5a, mRNA
	Homo sapiens adducin 2 (beta) (ADD2), transcript variant beta-3b, mRNA
NM_017484	Homo sapiens adducin 2 (beta) (ADD2), transcript variant beta-3a, mRNA
NM_017483	
NM_017482	Homo sapiens adducin 2 (beta) (ADD2), transcript variant beta-2, mRNA
NM_018561	Homo sapiens DKFZP586D2223 protein (DKFZP586D2223), mRNA
NM_018413	Homo sapiens chondroitin 4-sulfotransferase (C4ST), mRNA
NM_017835	Homo sapiens chromosome 21 open reading frame 59 (C21ORF59), mRNA
NM_018226	Homo sapiens arginyl aminopeptidase (aminopeptidase B)-like 1 (RNPEPL1),
ND 6 010004	mRNA
NM_018204	Homo sapiens cytoskeleton associated protein 2 (CKAP2), mRNA
NM_018200	Homo sapiens high-mobility group 20A (HMG20A), mRNA
NM_017595	Homo sapiens I-kappa-B-interacting Ras-like protein 2 (KBRAS2), mRNA
NM_017613	Homo sapiens downstream neighbor of SON (DONSON), mRNA
NM_017596	Homo sapiens KIAA0449 protein (KIAA0449), mRNA
NM_017456	Homo sapiens pleckstrin homology, Sec7 and coiled/coil domains 1(cytohesin 1) (PSCD1), transcript variant 2, mRNA
NM_016829	Homo sapiens 8-oxoguanine DNA glycosylase (OGG1), nuclear gene encoding
14141_010025	mitochondrial protein, transcript variant 2e, mRNA
NM_016828	Homo sapiens 8-oxoguanine DNA glycosylase (OGG1), nuclear gene encoding
_	mitochondrial protein, transcript variant 2d, mRNA
NM_016827	Homo sapiens 8-oxoguanine DNA glycosylase (OGG1), nuclear gene encoding
	mitochondrial protein, transcript variant 2c, mRNA
NM_016826	Homo sapiens 8-oxoguanine DNA glycosylase (OGG1), nuclear gene encoding mitochondrial protein, transcript variant 2b, mRNA
NM 016821	Homo sapiens 8-oxoguanine DNA glycosylase (OGG1), nuclear gene encoding
NWI_010821	mitochondrial protein, transcript variant 2a, mRNA
ND ( 01/020	Homo sapiens 8-oxoguanine DNA glycosylase (OGG1), nuclear gene encoding
NM_016820	mitochondrial protein, transcript variant 1c, mRNA
NM 016819	Homo sapiens 8-oxoguanine DNA glycosylase (OGG1), nuclear gene encoding
14141_010019	mitochondrial protein, transcript variant 1b, mRNA
NM 002197	Homo sapiens aconitase 1, soluble (ACO1), mRNA
NM 016841	Homo sapiens microtubule-associated protein tau (MAPT), transcript variant 4,
1	mRNA
NM 016835	Homo sapiens microtubule-associated protein tau (MAPT), transcript variant 1,
	mRNA
NM_016834	Homo sapiens microtubule-associated protein tau (MAPT), transcript variant 3,

	mRNA
NM 016938	Homo sapiens EGF-containing fibulin-like extracellular matrix protein 2
MMI_010339	(EFEMP2), mRNA
ND 4 005560	Homo sapiens LIM domain kinase 2 (LIMK2), transcript variant 2a, mRNA
NM_005569	Homo sapiens LIM domain kinase 2 (LIMK2), transcript variant 2b, mRNA
NM_016733	Homo sapiens LIM domain kinase 2 (LIMK1), transcript variant 1, mRNA
NM_002314	Homo sapiens LIM domain kinase 1 (LIMK1), transcript variant dLIMK, mRNA
NM_016735	Homo sapiens KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein
NM_006855	retention receptor 3 (KDELR3), transcript variant 1, mRNA
37 5 016657	Homo sapiens KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein
NM_016657	retention receptor 3 (KDELR3), transcript variant 2, mRNA
>D f 000101	Homo sapiens glycophorin C (Gerbich blood group) (GYPC), transcript variant
NM_002101	
27 f 01 C01 5	1, mRNA Homo sapiens glycophorin C (Gerbich blood group) (GYPC), transcript variant
NM_016815	2, mRNA
ND 4 005242	Homo sapiens coagulation factor II (thrombin) receptor-like 1 (F2RL1), mRNA
NM_005242	Homo sapiens ATP-binding cassette, sub-family G (WHITE), member 1
NM_016818	(ABCG1), transcript variant 2, mRNA
ND 6 004015	Homo sapiens ATP-binding cassette, sub-family G (WHITE), member 1
NM_004915	(ABCG1), transcript variant 1, mRNA
ND 4 002542	Homo sapiens 8-oxoguanine DNA glycosylase (OGG1), nuclear gene encoding
NM_002542	mitochondrial protein, transcript variant 1a, mRNA
ND 5 000/65	Homo sapiens acetylcholinesterase (YT blood group) (ACHE), transcript variant
NM_000665	E4-E6, mRNA
NT 6 012000	Homo sapiens mesenchyme homeo box 1 (MEOX1), transcript variant 2, mRNA
NM_013999	Homo sapiens methyl-CpG binding domain protein 2 (MBD2), transcript variant
NM_003927	1, mRNA
ND 6 015922	Homo sapiens methyl-CpG binding domain protein 2 (MBD2), transcript variant
NM_015832	testic specific mRNA
NM_002384	Homo sapiens methyl-CpG binding domain protein 1 (MBD1), transcript variant
1002364	1 mPNA
NM 015847	Homo sapiens methyl-CpG binding domain protein 1 (MBD1), transcript variant
1111_013647	PCM1 mRNA
NM 015846	Homo sapiens methyl-CpG binding domain protein 1 (MBD1), transcript variant
141VI_013640	1 mRNA
NM 015845	Homo sapiens methyl-CpG binding domain protein 1 (MBD1), transcript variant
14141_012642	12 mRNA
NM 015844	Homo sapiens methyl-CpG binding domain protein 1 (MBD1), transcript variant
14141_012044	3, mRNA
NM 002311	Homo sapiens ligase III, DNA, ATP-dependent (LIG3), transcript variant beta,
14141_002511	mRNA
NM_013975	Homo sapiens ligase III, DNA, ATP-dependent (LIG3), transcript variant alpha,
INIVI_0137/3	mRNA
NM 014190	Homo sapiens adducin 1 (alpha) (ADD1), transcript variant 3, mRNA
NM 014189	Homo sapiens adducin 1 (alpha) (ADD1), transcript variant 2, mRNA
NM 001119	Homo saniens adducin 1 (alpha) (ADD1), transcript variant 1, mRNA
	Homo sapiens acetylcholinesterase (YT blood group) (ACHE), transcript variant
NM_015831	E4-E5, mRNA
NIM 016572	Homo sapiens ubiquitin specific protease 21 (USP21), mRNA
NM_016572	Homo sapiens T-cell receptor interacting molecule (TRIM), mRNA
NM 016388	Homo sapiens transducer of ERBB2, 2 (TOB2), mRNA
NM 016272	Homo sapiens transcription factor ets (TEL2), mRNA
NM_016135	Home suprems transcription restor of (~===/2-====

NM_016247	Homo sapiens interphotoreceptor matrix proteoglycan 200 (SPACRCAN), mRNA
ND4 016224	
NM_016334	Homo sapiens putative G-protein coupled receptor (SH120), mRNA
NM_016124	Homo sapiens Rhesus blood group, D antigen (RHD), mRNA
NM_015865	Homo sapiens solute carrier family 14 (urea transporter), member 1 (Kidd blood group) (SLC14A1), mRNA
NM_016112	Homo sapiens polycystic kidney disease 2-like 1 (PKD2L1), mRNA
NM_016318	Homo sapiens purinergic receptor P2X, ligand-gated ion channel, 2 (P2RX2),
	mRNA
NM_016653	Homo sapiens sterile-alpha motif and leucine zipper containing kinase AZK
	(ZAK), mRNA
NM_016556	Homo sapiens GT198, complete ORF (HUMGT198A), mRNA
NM_016431	Homo sapiens mitogen-activated protein kinase 8 interacting protein 2
_	(MAPK8IP2), mRNA
NM_016377	Homo sapiens A kinase (PRKA) anchor protein 7 (AKAP7), mRNA
NM 016346	Homo sapiens nuclear receptor subfamily 2, group E, member 3 (NR2E3),
_	mRNA
NM 016325	Homo sapiens zinc finger protein 274 (ZNF274), mRNA
NM 016324	Homo sapiens zinc finger protein 274 (ZNF274), mRNA
NM 016293	Homo sapiens bridging integrator 2 (BIN2), mRNA
NM 015909	Homo sapiens neuroblastoma-amplified protein (LOC51594), mRNA
NM 015890	Homo sapiens spondyloepiphyseal dysplasia, late, pseudogene (SEDLP), mRNA
NM 015885	Homo sapiens PCF11p homolog (PCF11), mRNA
NM 015991	Homo sapiens complement component 1, q subcomponent, alpha polypeptide
	(C1QA), mRNA
NM 016201	Homo sapiens Leman coiled-coil protein (LCCP), mRNA
NM 016157	Homo sapiens trophinin (TRO), mRNA
NM 015869	Homo sapiens peroxisome proliferative activated receptor, gamma (PPARG),
_	mRNA
NM_016615	Homo sapiens solute carrier family 6 (neurotransmitter transporter, GABA),
	member 13 (SLC6A13), mRNA
NM_016389	Homo sapiens NS1-binding protein (NS1-BP), mRNA
NM_016648	Homo sapiens HDCMA18P protein (HDCMA18P), mRNA
NM_016527	Homo sapiens hydroxyacid oxidase 2 (long chain) (HAO2), mRNA
NM 016263	Homo sapiens Fzr1 protein (FZR1), mRNA
NM 016602	Homo sapiens G protein-coupled receptor 2 (GPR2), mRNA
NM 015892	Homo sapiens B cell RAG associated protein (BRAG), mRNA
NM_016187	Homo sapiens bridging integrator 2 (BIN2), mRNA
NM 003373	Homo sapiens vinculin (VCL), transcript variant VCL, mRNA
NM 014000	Homo sapiens vinculin (VCL), transcript variant meta-VCL, mRNA
NM_013992	Homo sapiens paired box gene 8 (PAX8), transcript variant PAX8E, mRNA
NM 013988	Homo sapiens Parkinson disease (autosomal recessive, juvenile) 2, parkin
_	(PARK2), transcript variant 3, mRNA
NM 013987	Homo sapiens Parkinson disease (autosomal recessive, juvenile) 2, parkin
	(PARK2), transcript variant 2, mRNA
NM 013985	Homo sapiens neuregulin 2 (NRG2), transcript variant 6, mRNA
NM 013984	Homo sapiens neuregulin 2 (NRG2), transcript variant 5, mRNA
NM_013983	Homo sapiens neuregulin 2 (NRG2), transcript variant 4, mRNA
NM 013982	Homo sapiens neuregulin 2 (NRG2), transcript variant 3, mRNA
NM_013981	Homo sapiens neuregulin 2 (NRG2), transcript variant 3, mRNA
NM 013964	Homo sapiens neuregulin 1 (NRG1), transcript variant HRG-alpha, mRNA
NM_013962	Homo sapiens neuregulin 1 (NRG1), transcript variant IRG-aipha, findva
010702	1

	CGE mPNA
NM_013961	Homo sapiens neuregulin 1 (NRG1), transcript variant GGF, mRNA
NM_013960	Homo sapiens neuregulin 1 (NRG1), transcript variant ndf43, mRNA
NM_013959	Homo sapiens neuregulin 1 (NRG1), transcript variant SMDF, mRNA
NM_013958	Homo sapiens neuregulin 1 (NRG1), transcript variant HRG-beta3, mRNA
NM_013957	Homo sapiens neuregulin 1 (NRG1), transcript variant HRG-beta2, mRNA
NM 013956	Homo sapiens neuregulin 1 (NRG1), transcript variant HRG-beta1, mRNA
NM 013955	Homo sapiens NADPH oxidase 1 (NOX1), transcript variant NOH-1Lv, mRNA
NM 013954	Homo sapiens NADPH oxidase 1 (NOX1), transcript variant NOH-1S, mRNA
NM_013995	Homo sapiens lysosomal-associated membrane protein 2 (LAMP2), transcript variant LAMP2B, mRNA
NM_007334	Homo sapiens killer cell lectin-like receptor subfamily D, member 1 (KLRD1),
NM_002262	Homo sapiens killer cell lectin-like receptor subfamily D, member 1 (KLRD1), transcript variant 1, mRNA
17.5 010076	Homo sapiens glutaryl-Coenzyme A dehydrogenase (GCDH), nuclear gene
NM_013976	encoding mitochondrial protein, transcript variant 2, mRNA
NR 6 01 50 41	Homo sapiens adenosine deaminase, RNA-specific (ADAR), transcript variant
NM_015841	ADAR-c, mRNA
ND 6 015940	Homo sapiens adenosine deaminase, RNA-specific (ADAR), transcript variant
NM_015840	ADAR h mŔNA
NM_001111	Homo sapiens adenosine deaminase, RNA-specific (ADAR), transcript variant
NM_001111	ADAR-a, mRNA
NM 014925	Homo sapiens KIAA1002 protein (KIAA1002), mRNA
NM 014905	Homo sapiens glutaminase (GLS), mRNA
NM 014833	Homo sapiens KIAA0618 gene product (KIAA0618), mRNA
NM 014863	Homo saniens B cell RAG associated protein (BRAG), mRNA
NM 015646	Homo conjens RAPIR member of RAS oncogene family (RAPIB), mRNA
NM_015423	Homo sapiens aminoadipate-semialdehyde dehydrogenase-phosphopantetheinyl
1411_013423	transferase (AASDHPPT), mRNA
NM_015523	Homo saniens small fragment nuclease (DKFZP566E144), mRNA
NM 014397	Homo saniens NIMA (never in mitosis gene a)-related kinase 6 (NEK6), mRNA
NM 014249	Homo sapiens nuclear receptor subfamily 2, group E, member 3 (NR2E3),
14141_014249	mRNA
NM 014361	Homo saniens contactin 5 (CNTN5), mRNA
NM 014341	Homo sapiens mitochondrial carrier homolog 1 (MTCH1), nuclear gene
11111_01.511	encoding mitochondrial protein, mRNA
NM 014556	Homo sapiens Ellis van Creveld syndrome (EVC), mRNA
NM_014306	Homo sapiens hypothetical protein (HSPC117), mRNA
NM 014593	Homo sapiens CpG binding protein (CGBP), mRNA
NM 014567	Homo saniens breast cancer anti-estrogen resistance 1 (BCAR1), mRNA
NM 014273	Homo sapiens a disintegrin-like and metalloprotease (reprolysin type) with
1417_01 1210	thrombospondin type 1 motif, 6 (ADAMTS6), mRNA
NM_014244	Homo saniens a disintegrin-like and metalloprotease (reprolysin type) with
11112_011211	thrombospondin type 1 motif, 2 (ADAMTS2), transcript variant 1, mRNA
NM 014449	Homo saniens protein A (A) transcript variant A-1, mRNA
NM_007319	
1111_00/515	374 mRNA
NM_007318	Homo sapiens presenilin 1 (Alzheimer disease 3) (PSEN1), transcript variant I-
NM 013953	Homo sapiens paired box gene 8 (PAX8), transcript variant PAX8D, mRNA
NM 013952	Homo saniens paired box gene 8 (PAX8), transcript variant PAX8C, mRNA
NM 013951	0 00 1 T 0 1 1 D A 7 0 D 20 A 1 A
141AT 012321	Tromo suprons pariod oor gone a version in

NM_013945	Homo sapiens paired box gene 7 (PAX7), transcript variant 2, mRNA
NM_013942	Homo sapiens paired box gene 3 (Waardenburg syndrome 1) (PAX3), transcript
	variant PAX3B, mRNA
NM_013411	Homo sapiens adenylate kinase 2 (AK2), nuclear gene encoding mitochondrial
	protein, transcript variant AK2B, mRNA
NM_000631	Homo sapiens neutrophil cytosolic factor 4 (40kD) (NCF4), transcript variant 1,
	mRNA
NM_013416	Homo sapiens neutrophil cytosolic factor 4 (40kD) (NCF4), transcript variant 2,
377	MRNA
NM_006125	Homo sapiens Rho GTPase activating protein 6 (ARHGAP6), transcript variant
37.5.010.10=	3, mRNA
NM_013427	Homo sapiens Rho GTPase activating protein 6 (ARHGAP6), transcript variant
3D 6 012402	1, mRNA
NM_013423	Homo sapiens Rho GTPase activating protein 6 (ARHGAP6), transcript variant
ND 4 012422	4, mRNA
NM_013422	Homo sapiens Rho GTPase activating protein 6 (ARHGAP6), transcript variant
NM 001174	5, mRNA
1001174	Homo sapiens Rho GTPase activating protein 6 (ARHGAP6), transcript variant
NM 013436	2, mRNA
NM 012310	Homo sapiens NCK-associated protein 1 (NCKAP1), mRNA
NM 013449	Homo sapiens kinesin family member 4A (KIF4A), mRNA
14141_015449	Homo sapiens bromodomain adjacent to zinc finger domain, 2A (BAZ2A), mRNA
NM_007333	Homo sapiens killer cell lectin-like receptor subfamily C, member 3 (KLRC3),
1111_007555	transcript variant NKG2-H, mRNA
NM_007328	Homo sapiens killer cell lectin-like receptor subfamily C, member 1 (KLRC1),
11.1.2.007520	transcript variant NKG2-B, mRNA
NM_002259	Homo sapiens killer cell lectin-like receptor subfamily C, member 1 (KLRC1),
_	transcript variant NKG2-A, mRNA
NM_004214	Homo sapiens fibroblast growth factor (acidic) intracellular binding protein
_	(FIBP), mRNA
NM_006350	Homo sapiens follistatin (FST), transcript variant FST317, mRNA
NM_013409	Homo sapiens follistatin (FST), transcript variant FST344, mRNA
NM_013324	Homo sapiens cytokine inducible SH2-containing protein (CISH), mRNA
NM_012486	Homo sapiens presenilin 2 (Alzheimer disease 4) (PSEN2), transcript variant 2,
-	mRNA
NM_012485	Homo sapiens hyaluronan-mediated motility receptor (RHAMM) (HMMR),
	transcript variant 2, mRNA
NM_012484	Homo sapiens hyaluronan-mediated motility receptor (RHAMM) (HMMR),
	transcript variant 1, mRNA
NM_012483	Homo sapiens granulysin (GNLY), transcript variant 519, mRNA
NM_006433	Homo sapiens granulysin (GNLY), transcript variant NKG5, mRNA
NM_001930	Homo sapiens deoxyhypusine synthase (DHPS), transcript variant 1, mRNA
NM_013407	Homo sapiens deoxyhypusine synthase (DHPS), transcript variant 3, mRNA
NM_013406	Homo sapiens deoxyhypusine synthase (DHPS), transcript variant 2, mRNA
NM_013229	Homo sapiens apoptotic protease activating factor (APAF1), transcript variant 1
) D ( 010000	mRNA
NM_013251	Homo sapiens tachykinin 3 (neuromedin K, neurokinin beta) (TAC3), mRNA
NM_013396	Homo sapiens ubiquitin specific protease 25 (USP25), mRNA
NM_013255	Homo sapiens muskelin 1, intracellular mediator containing kelch motifs
ND 4 012200	(MKLN1), mRNA
NM_013290	Homo sapiens GT198, complete ORF (HUMGT198A), mRNA

PCT/US03/05028 WO 03/074654

NM_005102	Homo sapiens fasciculation and elongation protein zeta 2 (zygin II) (FEZ2),
NM_004830	mRNA Homo sapiens cofactor required for Sp1 transcriptional activation, subunit 3
_	(130kD) (CRSP3), mRNA
NM_009588	Homo sapiens lymphotoxin beta (TNF superfamily, member 3) (LTB), transcript variant 2, mRNA
NM_013227	Homo sapiens aggrecan 1 (chondroitin sulfate proteoglycan 1, large aggregating proteoglycan, antigen identified by monoclonal antibody A0122) (AGC1), transcript variant 2, mRNA
NM_012475	Homo sapiens ubiquitin specific protease 21 (USP21), mRNA
NM_012428	Homo sapiens stromal cell derived factor receptor 1 (SDFR1), transcript variant beta mRNA
NM_012226	Homo sapiens purinergic receptor P2X, ligand-gated ion channel, 2 (P2RX2), mRNA
NM_012369	Homo sapiens olfactory receptor, family 2, subfamily F, member 1 (OR2F1), mRNA
NM_012218	Homo sapiens interleukin enhancer binding factor 3, 90kD (ILF3), mRNA
	Homo sapiens mitogen-activated protein kinase 8 interacting protein 2
NM_012324	(MAPK8IP2), mRNA
NM 012405	Homo sapiens isoprenylcysteine carboxyl methyltransferase (ICMT), mRNA
NM 012070	Homo sapiens attractin (ATRN), mRNA
NM 006874	Homo sapiens E74-like factor 2 (ets domain transcription factor) (ELF2), mRNA
	Homo sapiens synuclein, alpha (non A4 component of amyloid precursor)
NM_007308	(SNCA), transcript variant NACP112, mRNA
NM_000345	Homo sapiens synuclein, alpha (non A4 component of amyloid precursor) (SNCA), transcript variant NACP140, mRNA
NM 009589	Homo sapiens arylsulfatase D (ARSD), transcript variant 2, mRNA
NM_001158	Homo sapiens amine oxidase, copper containing 2 (retina-specific) (AOC2), transcript variant 1, mRNA
NM_005910	Homo sapiens microtubule-associated protein tau (MAPT), transcript variant 2, mRNA
NM_007338	Homo sapiens deleted in lung and esophageal cancer 1 (DLEC1), transcript variant DLEC1-L1, mRNA
NM_007337	Homo sapiens deleted in lung and esophageal cancer 1 (DLEC1), transcript variant DLEC1-S3, mRNA
NM_007336	Homo sapiens deleted in lung and esophageal cancer 1 (DLEC1), transcript variant DLEC1-S2, mRNA
NM_007335	Homo sapiens deleted in lung and esophageal cancer 1 (DLEC1), transcript variant DLEC1-S1, mRNA
NM_005106	Homo sapiens deleted in lung and esophageal cancer 1 (DLEC1), transcript variant DLEC1-N1, mRNA
NM_005002	Homo sapiens NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 9 (39kD) (NDUFA9), mRNA
NM 003771	Homo sapiens keratin, hair, acidic, 6 (KRTHA6), mRNA
NM_000438	Homo sapiens paired box gene 3 (Waardenburg syndrome 1) (PAX3), transcript variant PAX3A, mRNA
NM 007052	Homo sapiens NADPH oxidase 1 (NOX1), transcript variant NOH-1L, mRNA
NM 006715	Homo sapiens mannosidase, alpha, class 2C, member 1 (MAN2C1), mRNA
	Homo sapiens glutamate receptor, ionotrophic, AMPA 3 (GRIA3), transcript
NM_007325	veriant flin mRNA
NM_007325 NM_005813	variant flip, mRNA  Homo sapiens protein kinase C, nu (PRKCN), mRNA

	and disconsisted and di
	gene encoding mitochondrial protein, transcript variant M, mRNA
NM_007306	Homo sapiens breast cancer 1, early onset (BRCA1), transcript variant BRCA1-exon4, mRNA
NM_007305	Homo sapiens breast cancer 1, early onset (BRCA1), transcript variant BRCA1-delta9-10-11b, mRNA
NM_007304	Homo sapiens breast cancer 1, early onset (BRCA1), transcript variant BRCA1-delta11b, mRNA
NM_007303	Homo sapiens breast cancer 1, early onset (BRCA1), transcript variant BRCA1-delta11, mRNA
NM_007302	Homo sapiens breast cancer 1, early onset (BRCA1), transcript variant BRCA1-delta9-10, mRNA
NM_007301	Homo sapiens breast cancer 1, early onset (BRCA1), transcript variant BRCA1-
NM_007300	delta15-17, mRNA  Homo sapiens breast cancer 1, early onset (BRCA1), transcript variant BRCA1-
NM_007299	delta14-18, mRNA  Homo sapiens breast cancer 1, early onset (BRCA1), transcript variant BRCA1-
NM_007298	delta14-17, mRNA  Homo sapiens breast cancer 1, early onset (BRCA1), transcript variant BRCA1-
NM_007297	delta9-11, mRNA  Homo sapiens breast cancer 1, early onset (BRCA1), transcript variant BRCA1-delta2-10, mRNA
NM_007296	Homo sapiens breast cancer 1, early onset (BRCA1), transcript variant BRCA1a', mRNA
NM_007295	Homo sapiens breast cancer 1, early onset (BRCA1), transcript variant BRCA1b, mRNA
NM_007294	Homo sapiens breast cancer 1, early onset (BRCA1), transcript variant BRCA1a, mRNA
NM_007322	Homo sapiens RAN binding protein 3 (RANBP3), transcript variant RANBP3-d, mRNA
NM_007321	Homo sapiens RAN binding protein 3 (RANBP3), transcript variant RANBP3-c, mRNA
NM_007320	Homo sapiens RAN binding protein 3 (RANBP3), transcript variant RANBP3-b, mRNA
NM_000754	Homo sapiens catechol-O-methyltransferase (COMT), transcript variant MB-COMT, mRNA
NM_007310	Homo sapiens catechol-O-methyltransferase (COMT), transcript variant S-COMT, mRNA
NM_000714	Homo sapiens benzodiazapine receptor (peripheral) (BZRP), nuclear gene encoding mitochondrial protein, transcript variant PBR, mRNA
NM_007311	Homo sapiens benzodiazapine receptor (peripheral) (BZRP), nuclear gene encoding mitochondrial protein, transcript variant PBR-S, mRNA
NM_007314	Homo sapiens v-abl Abelson murine leukemia viral oncogene homolog 2 (arg, Abelson-related gene) (ABL2), transcript variant b, mRNA
NM_007313	Homo sapiens v-abl Abelson murine leukemia viral oncogene homolog 1 (ABL1), transcript variant b, mRNA
NM_005157	Homo sapiens v-abl Abelson murine leukemia viral oncogene homolog 1 (ABL1), transcript variant a, mRNA
NM 006325	Homo sapiens RAN, member RAS oncogene family (RAN), mRNA
NM_000902	Homo sapiens membrane metallo-endopeptidase (neutral endopeptidase, enkephalinase, CALLA, CD10) (MME), transcript variant 1, mRNA
NM_007289	Homo sapiens membrane metallo-endopeptidase (neutral endopeptidase, enkephalinase, CALLA, CD10) (MME), transcript variant 2b, mRNA

NM_007288	Homo sapiens membrane metallo-endopeptidase (neutral endopeptidase,
	enkephalinase, CALLA, CD10) (MME), transcript variant 2a, mRNA
NM_007287	Homo sapiens membrane metallo-endopeptidase (neutral endopeptidase,
_	enkephalinase, CALLA, CD10) (MME), transcript variant 1bis, mRNA
NM_006481	Homo sapiens transcription factor 2, hepatic; LF-B3; variant hepatic nuclear
_	factor (TCF2), transcript variant b, mRNA
NM_006884	Homo sapiens short stature homeobox 2 (SHOX2), transcript variant SHOX2a,
	mRNA
NM_003030	Homo sapiens short stature homeobox 2 (SHOX2), transcript variant SHOX2b,
1111_00000	mRNA
NM_003005	Homo sapiens selectin P (granule membrane protein 140kD, antigen CD62)
14141_003003	(SELP) mRNA
NM_006718	Homo sapiens pleiomorphic adenoma gene-like 1 (PLAGL1), transcript variant
NWI_000710	2, mRNA
NTM 005000	Homo sapiens solute carrier family 25 (mitochondrial carrier; phosphate carrier),
NM_005888	member 3 (SLC25A3), nuclear gene encoding mitochondrial protein, transcript
NR 006401	variant 1a, mRNA  Homo sapiens neuro-oncological ventral antigen 1 (NOVA1), transcript variant
NM_006491	
	3, mRNA Homo sapiens neuro-oncological ventral antigen 1 (NOVA1), transcript variant
NM_006489	
	2, mRNA
NM_007088	Homo sapiens calbindin 2, (29kD, calretinin) (CALB2), transcript variant
	CALB2c, mRNA
NM_007087	Homo sapiens calbindin 2, (29kD, calretinin) (CALB2), transcript variant
	CALB2b, mRNA
NM_001740	Homo sapiens calbindin 2, (29kD, calretinin) (CALB2), transcript variant
<del></del>	CALB2, mRNA
NM_007292	Homo sapiens acyl-Coenzyme A oxidase 1, palmitoyl (ACOX1), transcript
_	variant 2, mRNA
NM_004035	Homo sapiens acyl-Coenzyme A oxidase 1, palmitoyl (ACOX1), transcript
_	variant 1, mRNA
NM_000632	Homo sapiens integrin, alpha M (complement component receptor 3, alpha; also
	known as CD11b (p170), macrophage antigen alpha polypeptide) (ITGAM),
	mRNA
NM 007097	Homo sapiens clathrin, light polypeptide (Lcb) (CLTB), mRNA
NM 007099	Homo sapiens acid phosphatase 1, soluble (ACP1), transcript variant b, mRNA
	Homo sapiens TU3A protein (TU3A), mRNA
NM_007177 NM_007245	Homo sapiens ataxin 2 related protein (A2LP), transcript variant 1, mRNA
	Homo sapiens fibulin 1 (FBLN1), transcript variant A, mRNA
NM_006487	Homo sapiens fibulin 1 (FBLN1), transcript variant D, mRNA
NM_006486	Homo sapiens fibulin 1 (FBLN1), transcript variant B, mRNA  Homo sapiens fibulin 1 (FBLN1), transcript variant B, mRNA
NM_006485	HOMO Sapiens flouini i (FDLN1), nanscript variant D, matti
NM_006721	Homo sapiens adenosine kinase (ADK), transcript variant ADK-long, mRNA
NM_006132	Homo sapiens bone morphogenetic protein 1 (BMP1), transcript variant BMP1-
	4, mRNA
NM_006131	Homo sapiens bone morphogenetic protein 1 (BMP1), transcript variant BMP1-
	5, mRNA 1 (D) (D1) transported warriest PMP1
NM 006130	Homo sapiens bone morphogenetic protein 1 (BMP1), transcript variant BMP1-
	6 mRNA
NM_006129	Homo sapiens bone morphogenetic protein 1 (BMP1), transcript variant BMP1-
_	3 mRNA
NM_006128	- (D) (D) (
	2, mRNA

NM_002516	Homo sapiens neuro-oncological ventral antigen 2 (NOVA2), mRNA
NM_007008	Homo sapiens reticulon 4 (RTN4), mRNA
NM_007046	Homo sapiens elastin microfibril interface located protein (EMILIN), mRNA
NM_007037	Homo sapiens a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 8 (ADAMTS8), mRNA
NM_007038	Homo sapiens a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 5 (aggrecanase-2) (ADAMTS5), mRNA
NM 006799	Homo sapiens protease, serine, 21 (testisin) (PRSS21), mRNA
NM_006814	Homo sapiens proteasome (prosome, macropain) inhibitor subunit 1 (PI31) (PSMF1), mRNA
NM_003466	Homo sapiens paired box gene 8 (PAX8), transcript variant PAX8A, mRNA
NM_006790	Homo sapiens titin immunoglobulin domain protein (myotilin) (TTID), mRNA
NM 006782	Homo sapiens zinc finger protein-like 1 (ZFPL1), mRNA
NM_006795	Homo sapiens EH domain containing 1 (EHD1), mRNA
NM_006588	Homo sapiens sulfotransferase family, cytosolic, 1C, member 2 (SULT1C2), mRNA
NM_006694	Homo sapiens jumping translocation breakpoint (JTB), mRNA
NM_006597	Homo sapiens heat shock 70kD protein 8 (HSPA8), mRNA
NM_006708	Homo sapiens glyoxalase I (GLO1), mRNA
NM_006703	Homo sapiens nudix (nucleoside diphosphate linked moiety X)-type motif 3 (NUDT3), mRNA
NM_000655	Homo sapiens selectin L (lymphocyte adhesion molecule 1) (SELL), mRNA
NM_006488	Homo sapiens ketohexokinase (fructokinase) (KHK), transcript variant b, mRNA
NM_006297	Homo sapiens X-ray repair complementing defective repair in Chinese hamster cells 1 (XRCC1), mRNA
NM 006339	Homo sapiens high-mobility group 20B (HMG20B), mRNA
NM_006469	Homo sapiens NS1-binding protein (NS1-BP), mRNA
NM_006340	Homo sapiens BAI1-associated protein 2 (BAIAP2), transcript variant 3, mRNA
NM_001353	Homo sapiens aldo-keto reductase family 1, member C1 (dihydrodiol dehydrogenase 1; 20-alpha (3-alpha)-hydroxysteroid dehydrogenase) (AKR1C1), mRNA
NM_000202	Homo sapiens iduronate 2-sulfatase (Hunter syndrome) (IDS), transcript variant 1, mRNA
NM_005890	Homo sapiens growth arrest-specific 7 (GAS7), transcript variant b, mRNA
NM_006123	Homo sapiens iduronate 2-sulfatase (Hunter syndrome) (IDS), transcript variant 2, mRNA
NM_006053	Homo sapiens T-cell, immune regulator 1 (TCIRG1), mRNA
NM_005990	Homo sapiens serine/threonine kinase 10 (STK10), mRNA
NM_006019	Homo sapiens T-cell, immune regulator 1 (TCIRG1), mRNA
NM_006041	Homo sapiens heparan sulfate (glucosamine) 3-O-sulfotransferase 3B1 (HS3ST3B1), mRNA
NM_006042	Homo sapiens heparan sulfate (glucosamine) 3-O-sulfotransferase 3A1 (HS3ST3A1), mRNA
NM_006043	Homo sapiens heparan sulfate (glucosamine) 3-O-sulfotransferase 2 (HS3ST2), mRNA
NM_000557	Homo sapiens growth differentiation factor 5 (cartilage-derived morphogenetic protein-1) (GDF5), mRNA
NM_005847	Homo sapiens solute carrier family 23 (nucleobase transporters), member 2 (SLC23A2), mRNA
NM_005751	Homo sapiens A kinase (PRKA) anchor protein (yotiao) 9 (AKAP9), mRNA
NM_005691	Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 9 (ABCC9), transcript variant SUR2A, mRNA

	TD 1: 1: When the Complete Company of the Company o
NM_005688	Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 5 (ABCC5), mRNA
NM_005730	Homo sapiens conserved gene amplified in osteosarcoma (OS4), mRNA
NM 005562	Homo saniens laminin, gamma 2 (nicein (100kD), kalinin (105kD), BM600
	(100kD), Herlitz junctional epidermolysis bullosa)) (LAMC2), transcript variant
	1. mRNA
NM_005534	Homo sapiens interferon gamma receptor 2 (interferon gamma transducer 1)
11112_00000	(IFNGR2), mRNA
NM 005682	Homo sapiens G protein-coupled receptor 56 (GPR56), mRNA
NM_005666	Homo sapiens H factor (complement)-like 3 (HFL3), mRNA
NM 005503	Homo sapiens amyloid beta (A4) precursor protein-binding, family A, member 2
1111_003303	(X11-like) (APRA2) mRNA
NM_005431	Homo sapiens X-ray repair complementing defective repair in Chinese hamster
14141_002+21	cells 2 (XRCC2), mRNA
NM 005465	Homo sapiens v-akt murine thymoma viral oncogene homolog 3 (protein kinase
14141_005405	B, gamma) (AKT3), mRNA
NTM 005446	Homo sapiens purinergic receptor P2X-like 1, orphan receptor (P2RXL1),
NM_005446	mRNA
ND4 005226	Homo sapiens high density lipoprotein binding protein (vigilin) (HDLBP),
NM_005336	
37.5.005065	mRNA Homo sapiens gamma-glutamyltransferase 1 (GGT1), transcript variant 1,
NM_005265	
	mRNA Homo sapiens Ewing sarcoma breakpoint region 1 (EWSR1), transcript variant
NM_005243	
	EWS, mRNA
NM_005236	Homo sapiens excision repair cross-complementing rodent repair deficiency,
	complementation group 4 (ERCC4), mRNA
NM_005075	Homo sapiens solute carrier family 21 (organic anion transporter), member 3
	(SLC21A3), mRNA
NM_005050	Homo sapiens ATP-binding cassette, sub-family D (ALD), member 4 (ABCD4),
	transcript variant 1, mRNA
NM_005006	Homo sapiens NADH dehydrogenase (ubiquinone) Fe-S protein 1 (75kD)
	(NADH-coenzyme Q reductase) (NDUFS1), mRNA
NM_005135	Homo sapiens solute carrier family 12 (potassium/chloride transporters), member
	6 (SLC12A6), mRNA
NM_004968	Homo sapiens islet cell autoantigen 1 (69kD) (ICA1), transcript variant 2,
	mRNA (ICC 16 C 16 C 17 C 17 C 17 C 17 C 17 C 17
NM_005114	Homo sapiens heparan sulfate (glucosamine) 3-O-sulfotransferase 1 (HS3ST1),
	mRNA
NM_004958	Homo sapiens FK506 binding protein 12-rapamycin associated protein 1
	(FRAP1), mRNA
NM_001478	Homo sapiens UDP-N-acetyl-alpha-D-galactosamine:(N-acetylneuraminyl)-
	galactosylglucosylceramide N-acetylgalactosaminyltransferase (GalNAc-T)
	(GALGT), mRNA
NM_004031	Homo sapiens interferon regulatory factor 7 (IRF7), transcript variant d, mRNA
NM_004030	Homo sapiens interferon regulatory factor 7 (IRF7), transcript variant c, mRNA
NM_004029	Homo sapiens interferon regulatory factor 7 (IRF7), transcript variant b, mRNA
NM 004034	Homo sapiens annexin A7 (ANXA7), transcript variant 2, mRNA
NM 001156	Homo sapiens annexin A7 (ANXA7), transcript variant 1, mRNA
NM 004033	Homo sapiens annexin A6 (ANXA6), transcript variant 2, mRNA
NM 001155	Homo sapiens annexin A6 (ANXA6), transcript variant 1, mRNA
NM 004629	Homo sapiens Fanconi anemia, complementation group G (FANCG), mRNA
	Homo sapiens VAMP (vesicle-associated membrane protein)-associated protein
NM_004738_	Homo sapiens VAMP (vesicle-associated memorane protein)-associated protein

	D. IC(VADD)DNA
ND 6 004774	B and C (VAPB), mRNA
NM_004774	Homo sapiens PPAR binding protein (PPARBP), mRNA
NM_004819	Homo sapiens symplekin; Huntingtin interacting protein I (SPK), mRNA
NM_004169	Homo sapiens serine hydroxymethyltransferase 1 (soluble) (SHMT1), mRNA
NM_004186	Homo sapiens sema domain, immunoglobulin domain (Ig), short basic domain,
	secreted, (semaphorin) 3F (SEMA3F), mRNA
NM_004730	Homo sapiens eukaryotic translation termination factor 1 (ETF1), mRNA
NM_004161	Homo sapiens RAB1, member RAS oncogene family (RAB1), mRNA
NM_004762	Homo sapiens pleckstrin homology, Sec7 and coiled/coil domains 1(cytohesin 1) (PSCD1), transcript variant 1, mRNA
NM 004253	Homo sapiens phospholipase A2-activating protein (PLAA), mRNA
NM_004562	Homo sapiens Parkinson disease (autosomal recessive, juvenile) 2, parkin (PARK2), transcript variant 1, mRNA
NM_004705	Homo sapiens protein-kinase, interferon-inducible double stranded RNA
11112_00 1100	dependent inhibitor, repressor of (P58 repressor) (PRKRIR), mRNA
NM_004883	Homo sapiens neuregulin 2 (NRG2), transcript variant 1, mRNA
NM 004559	Homo sapiens nuclease sensitive element binding protein 1 (NSEP1), mRNA
NM 004646	Homo sapiens nephrosis 1, congenital, Finnish type (nephrin) (NPHS1), mRNA
NM 004897	Homo sapiens multiple inositol polyphosphate phosphatase 1 (MINPP1), mRNA
NM 004527	Homo sapiens mesenchyme homeo box 1 (MEOX1), transcript variant 1, mRNA
NM 004912	Homo sapiens cerebral cavernous malformations 1 (CCM1), mRNA
NM 001572	Homo sapiens interferon regulatory factor 7 (IRF7), transcript variant a, mRNA
NM 004516	Homo sapiens interleukin enhancer binding factor 3, 90kD (ILF3), mRNA
NM 004505	Homo sapiens ubiquitin specific protease 6 (Tre-2 oncogene) (USP6), mRNA
	Homo sapiens RAB2, member RAS oncogene family-like (RAB2L), mRNA
NM_004761	
NM_004495	Homo sapiens neuregulin 1 (NRG1), transcript variant HRG-gamma, mRNA
NM_004821	Homo sapiens heart and neural crest derivatives expressed 1 (HAND1), mRNA
NM_004458	Homo sapiens fatty-acid-Coenzyme A ligase, long-chain 4 (FACL4), transcript variant 1, mRNA
NM_004091	Homo sapiens E2F transcription factor 2 (E2F2), mRNA
NM_004714	Homo sapiens dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 1B (DYRK1B), transcript variant a, mRNA
ND ( 004950	
NM_004859	Homo sapiens clathrin, heavy polypeptide (Hc) (CLTC), mRNA
NM_004921	Homo sapiens chloride channel, calcium activated, family member 3 (CLCA3), mRNA
NM_004344	Homo sapiens centrin, EF-hand protein, 2 (CETN2), mRNA
NM_004332	Homo sapiens biphenyl hydrolase-like (serine hydrolase; breast epithelial mucin-associated antigen) (BPHL), mRNA
NM_004842	Homo sapiens A kinase (PRKA) anchor protein 7 (AKAP7), mRNA
NM_004194	Homo sapiens a disintegrin and metalloproteinase domain 22 (ADAM22), mRNA
NM_004300	Homo sapiens acid phosphatase 1, soluble (ACP1), transcript variant a, mRNA
NM_004769	Homo sapiens amiloride-sensitive cation channel 3, testis (ACCN3), transcript variant 1, mRNA
NM_004027	Homo sapiens inositol polyphosphate-4-phosphatase, type I, 107kD (INPP4A), transcript variant a, mRNA
NM 004003	Homo sapiens carnitine acetyltransferase (CRAT), nuclear gene encoding
	mitochondrial protein, transcript variant peroxisomal, mRNA
NM_004028	Homo sapiens aquaporin 4 (AQP4), transcript variant b, mRNA
NM_001650	Homo sapiens aquaporin 4 (AQP4), transcript variant a, mRNA
NM_002390	Homo sapiens a disintegrin and metalloproteinase domain 11 (ADAM11),
L	transcript variant 1, mRNA

	TO A STATE OF THE PARTY OF THE
NM_001604	Homo sapiens paired box gene 6 (aniridia, keratitis) (PAX6), mRNA
NM_003995	Homo sapiens natriuretic peptide receptor B/guanylate cyclase B (atrionatriuretic
	peptide receptor B) (NPR2), mRNA
NM_003994	Homo sapiens KIT ligand (KITLG), mRNA
NM_001063	Homo sapiens transferrin (TF), mRNA
NM 003990	Homo sapiens paired box gene 2 (PAX2), transcript variant e, mRNA
NM 003989	Homo sapiens paired box gene 2 (PAX2), transcript variant d, mRNA
NM 003988	Homo sapiens paired box gene 2 (PAX2), transcript variant c, mRNA
NM 003987	Homo sapiens paired box gene 2 (PAX2), transcript variant a, mRNA
NM 000278	Homo sapiens paired box gene 2 (PAX2), transcript variant b, mRNA
NM 000278	Homo sapiens ketohexokinase (fructokinase) (KHK), transcript variant a, mRNA
NM 000115	Homo sapiens endothelin receptor type B (EDNRB), transcript variant 1, mRNA
	Homo sapiens carnitine acetyltransferase (CRAT), nuclear gene encoding
NM_000755	mitochondrial protein, transcript variant mitochondrial, mRNA
	mitochondriai protein, transcript variant interestript variant pholicity mRNA
NM_001292	Homo sapiens CDC-like kinase 3 (CLK3), transcript variant phelk3/152, mRNA
NM_001291	Homo sapiens CDC-like kinase 2 (CLK2), transcript variant phclk2/139, mRNA
NM_001282	Homo sapiens adaptor-related protein complex 2, beta 1 subunit (AP2B1), mRNA
NM 001272	Homo sapiens chromodomain helicase DNA binding protein 3 (CHD3), mRNA
NM 001268	Homo sapiens chromosome condensation 1-like (CHC1L), mRNA
	Homo sapiens CD3Z antigen, zeta polypeptide (TiT3 complex) (CD3Z), mRNA
NM_000734	Homo sapiens B-cell CLL/lymphoma 2 (BCL2), nuclear gene encoding
NM_000657	mitochondrial protein, transcript variant beta, mRNA
15.000.000	mitochondrial protein, transcript variant octa, micros
NM_000633	Homo sapiens B-cell CLL/lymphoma 2 (BCL2), nuclear gene encoding
	mitochondrial protein, transcript variant alpha, mRNA
NM_000055	Homo sapiens butyrylcholinesterase (BCHE), mRNA
NM_003594	Homo sapiens transcription termination factor, RNA polymerase II (TTF2), mRNA
NM_003722	Homo sapiens tumor protein 63 kDa with strong homology to p53 (TP63), mRNA
NM 003856	Homo sapiens interleukin 1 receptor-like 1 (IL1RL1), mRNA
NM 003140	Homo sapiens sex determining region Y (SRY), mRNA
NM_003615	Homo sapiens solute carrier family 4, sodium bicarbonate cotransporter, member
14141_005015	7 (SLC4A7), mRNA
NM_003759	Homo sapiens solute carrier family 4, sodium bicarbonate cotransporter, member
	4 (SLC4A4), mRNA
NM_002980	Homo sapiens secretin receptor (SCTR), mRNA
NM_002890	Homo sapiens RAS p21 protein activator (GTPase activating protein) 1 (RASA1), transcript variant 1, mRNA
NM_003624	Homo sapiens RAN binding protein 3 (RANBP3), transcript variant RANBP3-a, mRNA
NM_002817	Homo sapiens proteasome (prosome, macropain) 26S subunit, non-ATPase, 13
_	(PSMD13), mRNA
NM_000447	Homo sapiens presenilin 2 (Alzheimer disease 4) (PSEN2), transcript variant 1, mRNA
NM_000021	Homo sapiens presenilin 1 (Alzheimer disease 3) (PSEN1), transcript variant I-
	467, mRNA
NM_002768	Homo sapiens procollagen (type III) N-endopeptidase (PCOLN3), mRNA
NM_002752	Homo sapiens mitogen-activated protein kinase 9 (MAPK9), mRNA
NM_002656	Homo sapiens pleiomorphic adenoma gene-like 1 (PLAGL1), transcript variant 1, mRNA
NM 002635	Homo sapiens solute carrier family 25 (mitochondrial carrier; phosphate carrier),
11111	

	1 - 2 (CI C25 A2) los disc ita chandrial
	member 3 (SLC25A3), nuclear gene encoding mitochondrial protein, transcript variant 1b, mRNA
NM_002584	Homo sapiens paired box gene 7 (PAX7), transcript variant 1, mRNA
NM_000280	Homo sapiens paired box gene 6 (aniridia, keratitis) (PAX6), mRNA
NM_002555	Homo sapiens solute carrier family 22 (organic cation transporter), member 1-like (SLC22A1L), mRNA
NM_000907	Homo sapiens natriuretic peptide receptor B/guanylate cyclase B (atrionatriuretic peptide receptor B) (NPR2), mRNA
NM_002515	Homo sapiens neuro-oncological ventral antigen 1 (NOVA1), transcript variant 1, mRNA
NM 003204	Homo sapiens nuclear factor (erythroid-derived 2)-like 1 (NFE2L1), mRNA
NM 003970	Homo sapiens myomesin (M-protein) 2 (165kD) (MYOM2), mRNA
NM 000899	Homo sapiens KIT ligand (KITLG), mRNA
NM_002394	Homo sapiens solute carrier family 3 (activators of dibasic and neutral amino acid transport), member 2 (SLC3A2), mRNA
NM_001879	Homo sapiens mannan-binding lectin serine protease 1 (C4/C2 activating component of Ra-reactive factor) (MASP1), mRNA
NM_002353	Homo sapiens tumor-associated calcium signal transducer 2 (TACSTD2), mRNA
NM_002341	Homo sapiens lymphotoxin beta (TNF superfamily, member 3) (LTB), transcript variant 1, mRNA
NM_002294	Homo sapiens lysosomal-associated membrane protein 2 (LAMP2), transcript variant LAMP2A, mRNA
NM 002264	Homo sapiens karyopherin alpha 1 (importin alpha 5) (KPNA1), mRNA
NM_002261	Homo sapiens killer cell lectin-like receptor subfamily C, member 3 (KLRC3), transcript variant NKG2-E, mRNA
NM 002230	Homo sapiens junction plakoglobin (JUP), transcript variant 1, mRNA
NM_001566	Homo sapiens inositol polyphosphate-4-phosphatase, type I, 107kD (INPP4A), transcript variant b, mRNA
NM 002164	Homo sapiens indoleamine-pyrrole 2,3 dioxygenase (INDO), mRNA
NM_003822	Homo sapiens nuclear receptor subfamily 5, group A, member 2 (NR5A2), mRNA
NM_000836	Homo sapiens glutamate receptor, ionotropic, N-methyl D-aspartate 2D (GRIN2D), mRNA
NM_000828	Homo sapiens glutamate receptor, ionotrophic, AMPA 3 (GRIA3), transcript variant flop, mRNA
NM 002056	Homo sapiens glutamine-fructose-6-phosphate transaminase 1 (GFPT1), mRNA
NM_000161	Homo sapiens GTP cyclohydrolase 1 (dopa-responsive dystonia) (GCH1), mRNA
NM_000159	Homo sapiens glutaryl-Coenzyme A dehydrogenase (GCDH), nuclear gene encoding mitochondrial protein, transcript variant 1, mRNA
NM 003644	Homo sapiens growth arrest-specific 7 (GAS7), transcript variant a, mRNA
NM_000817	Homo sapiens glutamate decarboxylase 1 (brain, 67kD) (GAD1), transcript variant GAD67, mRNA
NM_000813	Homo sapiens gamma-aminobutyric acid (GABA) A receptor, beta 2 (GABRB2), transcript variant 2, mRNA
NM_000146	Homo sapiens ferritin, light polypeptide (FTL), mRNA
NM_001996	Homo sapiens fibulin 1 (FBLN1), transcript variant C, mRNA
NM_001995	Homo sapiens fatty-acid-Coenzyme A ligase, long-chain 1 (FACL1), nuclear gene encoding mitochondrial protein, mRNA
NM_001973	Homo sapiens ELK4, ETS-domain protein (SRF accessory protein 1) (ELK4), transcript variant a, mRNA

	D (EDNDD) transcript variant 2 mRNA
NM_003991	Homo sapiens endothelin receptor type B (EDNRB), transcript variant 2, mRNA
NM_001925	Homo sapiens defensin, alpha 4, corticostatin (DEFA4), mRNA
NM_001359	Homo sapiens 2,4-dienoyl CoA reductase 1, mitochondrial (DECR1), nuclear
	gene encoding mitochondrial protein, mRNA
NM_001337	Homo sapiens chemokine (C-X3-C) receptor 1 (CX3CR1), mRNA
NM_001835	Homo sapiens clathrin, heavy polypeptide-like 1 (CLTCL1), transcript variant 1, mRNA
NTM ( 001924	Homo sapiens clathrin, light polypeptide (Lcb) (CLTB), transcript variant
NM_001834	nonbrain, mRNA
ND ( 002002	Homo sapiens CDC-like kinase 3 (CLK3), transcript variant phelk3, mRNA
NM_003992	Homo sapiens CDC-like kinase 2 (CLK2), transcript variant phelk2, mRNA
NM_003993	Homo sapiens chloride channel 6 (CLCN6), transcript variant ClC-6a, mRNA
NM_001286	Homo sapiens chloride channel, calcium activated, family member 1 (CLCA1),
NM_001285	
37.6 001005	mRNA Homo sapiens creatine kinase, mitochondrial 2 (sarcomeric) (CKMT2), nuclear
NM_001825	Homo sapiens creatine kinase, introchondriai 2 (salconterio) (Creating), national
	gene encoding mitochondrial protein, mRNA
NM_003465	Homo sapiens chitinase 1 (chitotriosidase) (CHIT1), mRNA
NM_001783	Homo sapiens CD79A antigen (immunoglobulin-associated alpha) (CD79A),
	transcript variant 1, mRNA  Homo sapiens bone morphogenetic protein 1 (BMP1), transcript variant BMP1-
NM_001199	
	1, mRNA
NM_001669	Homo sapiens arylsulfatase D (ARSD), transcript variant 1, mRNA
NM_001170	Homo sapiens aquaporin 7 (AQP7), mRNA
NM_001160	Homo sapiens apoptotic protease activating factor (APAF1), transcript variant 2, mRNA
NM 001149	Homo sapiens ankyrin 3, node of Ranvier (ankyrin G) (ANK3), transcript variant
_	2. mRNA
NM_001625	Homo sapiens adenylate kinase 2 (AK2), nuclear gene encoding mitochondrial
	protein, transcript variant AK2A, mRNA
NM_001135	Homo sapiens aggrecan 1 (chondroitin sulfate proteoglycan 1, large aggregating
	proteoglycan, antigen identified by monoclonal antibody A0122) (AGC1),
	transcript variant 1, mRNA
NM_001123	Homo sapiens adenosine kinase (ADK), transcript variant ADK-short, mRNA
NM_003812	Homo sapiens a disintegrin and metalloproteinase domain 23 (ADAM23), mRNA
NM 001095	Homo sapiens amiloride-sensitive cation channel 2, neuronal (ACCN2),
14141_001055	transcript variant 2, mRNA
NM 016184	Homo sapiens C-type (calcium dependent, carbohydrate-recognition domain)
14141_010104	lectin, superfamily member 6 (CLECSF6), mRNA
NM 003186	Homo sapiens transgelin (TAGLN), mRNA
NM 004084	Homo sapiens defensin, alpha 1, myeloid-related sequence (DEFA1), mRNA
NM 022908	Homo sapiens hypothetical protein FLJ12442 (FLJ12442), mRNA
	Homo sapiens hypothetical protein FLJ13195 similar to stromal antigen 3
NM_022906	(FLJ13195), mRNA
NM 022903	Homo sapiens hypothetical protein FLJ12800 (FLJ12800), mRNA
NM 022902	Homo sapiens hypothetical protein FLJ12496 (FLJ12496), mRNA
NM 022902	Homo sapiens hypothetical protein FLJ21213 (FLJ21213), mRNA
NM 022895	Homo sapiens hypothetical protein FLJ12448 (FLJ12448), mRNA
	Homo sapiens transforming, acidic coiled-coil containing protein 2 (TACC2),
NM_006997	mRNA
NM 020979	Homo sapiens adaptor protein with pleckstrin homology and src homology 2
1	domains (APS), mRNA

NM 018557	Homo sapiens low density lipoprotein-related protein 1B (deleted in tumors)
11111_010337	(LRP1B), mRNA
NM 014921	Homo sapiens lectomedin-2 (KIAA0821), mRNA
NM 014112	Homo sapiens trichorhinophalangeal syndrome I gene (TRPS1), mRNA
NM_000539	Homo sapiens rhodopsin (opsin 2, rod pigment) (retinitis pigmentosa 4,
	autosomal dominant) (RHO), mRNA
NM_012452	Homo sapiens transmembrane activator and CAML interactor (TACI), mRNA
NM_003564	Homo sapiens transgelin 2 (TAGLN2), mRNA
NM_003632	Homo sapiens contactin associated protein 1 (CNTNAP1), mRNA
NM_006506	Homo sapiens RAS p21 protein activator 2 (RASA2), mRNA
NM_014427	Homo sapiens copine VII (CPNE7), mRNA
NM_006032	Homo sapiens copine VI (neuronal) (CPNE6), mRNA
NM_005338	Homo sapiens huntingtin interacting protein 1 (HIP1), mRNA
NM_021973	Homo sapiens heart and neural crest derivatives expressed 2 (HAND2), mRNA
NM_005339	Homo sapiens huntingtin interacting protein 2 (HIP2), mRNA
NM_021920	Homo sapiens secretin (SCT), mRNA
NM_016491	Homo sapiens mitochondrial ribosomal protein L37 (MRPL37), mRNA
NM_014211	Homo sapiens gamma-aminobutyric acid (GABA) A receptor, pi (GABRP), mRNA
NM_004658	Homo sapiens RAS protein activator like 1 (GAP1 like) (RASAL1), mRNA
NM 004807	Homo sapiens heparan sulfate 6-O-sulfotransferase (HS6ST), mRNA
NM_002622	Homo sapiens prefoldin 1 (PFDN1), mRNA
NM 005186	Homo sapiens calpain 1, (mu/I) large subunit (CAPN1), mRNA
NM 001748	Homo sapiens calpain 2, (m/II) large subunit (CAPN2), mRNA
NM_014299	Homo sapiens bromodomain-containing 4 (BRD4), mRNA
NM_007208	Homo sapiens mitochondrial ribosomal protein L3 (MRPL3), mRNA
NM_022838	Homo sapiens hypothetical protein FLJ12969 (FLJ12969), mRNA
NM_022837	Homo sapiens hypothetical protein FLJ22833 (FLJ22833), mRNA
NM_022830	Homo sapiens hypothetical protein FLJ22347 (FLJ22347), mRNA
NM_022819	Homo sapiens phospholipase A2, group IIF (PLA2G2F), mRNA
NM_020245	Homo sapiens tubby super-family protein (TUSP), mRNA
NM_020061	Homo sapiens opsin 1 (cone pigments), long-wave-sensitive (color blindness, protan) (OPN1LW), mRNA
NM_000513	Homo sapiens opsin 1 (cone pigments), medium-wave-sensitive (color blindness, deutan) (OPN1MW), mRNA
NM 001708	Homo sapiens opsin 1 (cone pigments), short-wave-sensitive (color blindness,
1441_001700	tritan) (OPN1SW), mRNA
NM 016363	Homo sapiens glycoprotein VI (platelet) (GP6), mRNA
NM_022139	Homo sapiens GDNF family receptor alpha 4 (GFRA4), mRNA
NM 002485	Homo sapiens Nijmegen breakage syndrome 1 (nibrin) (NBS1), mRNA
NM 006052	Homo sapiens Down syndrome critical region gene 3 (DSCR3), mRNA
NM 005867	Homo sapiens Down syndrome critical region gene 4 (DSCR4), mRNA
NM 005087	Homo sapiens fragile X mental retardation, autosomal homolog 1 (FXR1),
_	mRNA
NM 004403	Homo sapiens deafness, autosomal dominant 5 (DFNA5), mRNA
NM_000433	Homo sapiens neutrophil cytosolic factor 2 (65kD, chronic granulomatous
-	disease, autosomal 2) (NCF2), mRNA
NM_000111	Homo sapiens solute carrier family 26, member 3 (SLC26A3), mRNA
NM_000044	Homo sapiens androgen receptor (dihydrotestosterone receptor; testicular
_	feminization; spinal and bulbar muscular atrophy; Kennedy disease) (AR), mRNA
NM 000333	Homo sapiens spinocerebellar ataxia 7 (olivopontocerebellar atrophy with retinal
TATAT 000222	Tromo sapiens spinocerebenar ataxia / (onvopontocerebenar atrophy with retinar

	degeneration) (SCA7), mRNA
NM_003776	Homo sapiens nuclear localization signal deleted in velocardiofacial syndrome
	(NI.VCF), mRNA
NM 003941	Homo sapiens Wiskott-Aldrich syndrome-like (WASL), mRNA
NM 020680	Homo sapiens N-terminal kinase-like (NTKL), mRNA
NM 022789	Homo senions interleukin 17F (II.17F), mRNA
NM_022787	Homo sapiens NMN adenylyltransferase; nicotinamide mononucleotide adenylyl transferase (NMNAT), mRNA
T 6 000786	Homo sapiens likely ortholog of yeast ARV1 (ARV1), mRNA
NM_022786	Homo sapiens hypothetical protein FLJ23588 (FLJ23588), mRNA
NM 022785	Homo sapiens hypothetical protein FLJ22127 (FLJ22127), mRNA
NM_022775	Homo sapiens hypothetical protein FLJ12681 (FLJ12681), mRNA
NM_022773	Homo sapiens hypothetical protein FL312001 (FL312001), mRNA
NM_022772	Homo sapiens hypothetical protein FLJ21935 (FLJ21935), mRNA
NM_022761	Homo sapiens hypothetical protein FLJ23499 (FLJ23499), mRNA
NM_022756	Homo sapiens hypothetical protein FLJ11730 (FLJ11730), mRNA
NM_022739	Homo sapiens E3 ubiquitin ligase SMURF2 (SMURF2), mRNA
NM_022725	Homo sapiens Fanconi anemia, complementation group F (FANCF), mRNA
NM_017646	Homo sapiens tRNA isopentenylpyrophosphate transferase (IPT), mRNA
NM_005443	Homo sapiens 3'-phosphoadenosine 5'-phosphosulfate synthase 1 (PAPSS1), mRNA
NM_004670	Homo sapiens 3'-phosphoadenosine 5'-phosphosulfate synthase 2 (PAPSS2),
NM_001084	Homo sapiens procollagen-lysine, 2-oxoglutarate 5-dioxygenase 3 (PLOD3),
14141_001004	mPNA
NM 022720	Homo saniens DiGeorge syndrome critical region gene 8 (DGCR8), mRNA
NM_007331	Homo sapiens Wolf-Hirschhorn syndrome candidate 1 (WHSC1), mRNA
NM_007123	Homo sapiens Usher syndrome 2A (autosomal recessive, mild) (USH2A),
_	mRNA
NM 000553	Homo sapiens Werner syndrome (WRN), mRNA
NM_006531	Homo sapiens Probe hTg737 (polycystic kidney disease, autosomal recessive, in) (TG737), mRNA
NM 018962	Homo sapiens Down syndrome critical region gene 6 (DSCR6), mRNA
NM 018848	Homo sapiens McKusick-Kaufman syndrome (MKKS), mRNA
NM_017424	Homo sapiens cat eye syndrome chromosome region, candidate I (CECRI),
DD 6 01 5000	mRNA Homo sapiens TPA inducible gene-1 (TIG-1), mRNA
NM_015889	Homo sapiens TPA inducible gene-1 (TIG-1), inicutal Homo sapiens Down syndrome critical region gene 5 (DSCR5), mRNA
NM_016430	Homo sapiens Down syndrome critical region gene 1 (DSCR1), mRNA  Homo sapiens Down syndrome critical region gene 1 (DSCR1), mRNA
NM_004414	Homo sapiens Down syndrome critical region gene 1-like 2 (DSCR1L2), mRNA  Homo sapiens Down syndrome critical region gene 1-like 2 (DSCR1L2), mRNA
NM_013441	Homo sapiens Down syndrome critical region gene 1-ince 2 (Docted), inclusion
NM_012436	Homo sapiens sperm associated antigen 8 (SPAG8), mRNA
NM_012227	Homo sapiens Pseudoautosomal GTP-binding protein-like (PGPL), mRNA
NM_007173	Homo sapiens protease, serine, 23 (SPUVE), mRNA
NM_000501	Homo sapiens elastin (supravalvular aortic stenosis, Williams-Beuren syndrome (ELN), mRNA
NM 006025	Homo sapiens protease, serine, 22 (P11), mRNA
NM_005609	Homo sapiens phosphorylase, glycogen; muscle (McArdle syndrome, glycogen storage disease type V) (PYGM), mRNA
NM_004991	Homo saniens myelodysplasia syndrome 1 (MDS1), mRNA
NM_004600	Homo sapiens Sjogren syndrome antigen A2 (60kD, ribonucleoprotein
NM_004380	Homo sapiens CREB binding protein (Rubinstein-Taybi syndrome) (CREBBP),

NM_000551	Homo sapiens von Hippel-Lindau syndrome (VHL), mRNA
NM_000462	Homo sapiens ubiquitin protein ligase E3A (human papilloma virus E6-
_	associated protein, Angelman syndrome) (UBE3A), mRNA
NM 001064	Homo sapiens transketolase (Wernicke-Korsakoff syndrome) (TKT), mRNA
NM 000356	Homo sapiens Treacher Collins-Franceschetti syndrome 1 (TCOF1), mRNA
NM 000455	Homo sapiens serine/threonine kinase 11 (Peutz-Jeghers syndrome) (STK11),
	mRNA
NM 002351	Homo sapiens SH2 domain protein 1A, Duncan's disease (lymphoproliferative
	syndrome) (SH2D1A), mRNA
NM 000336	Homo sapiens sodium channel, nonvoltage-gated 1, beta (Liddle syndrome)
1111_000550	(SCNN1B), mRNA
NM_000335	Homo sapiens sodium channel, voltage-gated, type V, alpha polypeptide (long
11111_000333	(electrocardiographic) QT syndrome 3) (SCN5A), mRNA
NM_000318	Homo sapiens peroxisomal membrane protein 3 (35kD, Zellweger syndrome)
14141_000518	(PXMP3), mRNA
NM_000311	Homo sapiens prion protein (p27-30) (Creutzfeld-Jakob disease, Gerstmann-
14141_000511	Strausler-Scheinker syndrome, fatal familial insomnia) (PRNP), mRNA
NM_000299	Homo sapiens plakophilin 1 (ectodermal dysplasia/skin fragility syndrome)
NWI_000299	(PKP1), mRNA
NM 000283	
NWI_000283	Homo sapiens phosphodiesterase 6B, cGMP-specific, rod, beta (congenital
NTM 002721	stationary night blindness 3, autosomal dominant) (PDE6B), mRNA
NM_003731	Homo sapiens Sjogren's syndrome nuclear autoantigen 1 (SSNA1), mRNA
NM_000260	Homo sapiens myosin VIIA (Usher syndrome 1B (autosomal recessive, severe))
NR 6 0007700	(MYO7A), mRNA
NM_003720	Homo sapiens Down syndrome critical region gene 2 (DSCR2), mRNA
NM_000195	Homo sapiens Hermansky-Pudlak syndrome (HPS), mRNA
NM_000194	Homo sapiens hypoxanthine phosphoribosyltransferase 1 (Lesch-Nyhan
	syndrome) (HPRT1), mRNA
NM_000171	Homo sapiens glycine receptor, alpha 1 (startle disease/hyperekplexia, stiff man
	syndrome) (GLRA1), mRNA
NM_003494	Homo sapiens dysferlin, limb girdle muscular dystrophy 2B (autosomal
	recessive) (DYSF), mRNA
NM_000081	Homo sapiens Chediak-Higashi syndrome 1 (CHS1), mRNA
NM_000052	Homo sapiens ATPase, Cu++ transporting, alpha polypeptide (Menkes
	syndrome) (ATP7A), mRNA
NM_001635	Homo sapiens amphiphysin (Stiff-Mann syndrome with breast cancer 128kD
	autoantigen) (AMPH), mRNA
NM_022663	Homo sapiens CTAGE-1 protein (CTAGE-1), mRNA
NM_022662	Homo sapiens meiotic checkpoint regulator (MCPR), mRNA
NM_022658	Homo sapiens homeo box C8 (HOXC8), mRNA
NM_000569	Homo sapiens Fc fragment of IgG, low affinity IIIa, receptor for (CD16)
	(FCGR3A), mRNA
NM_000802	Homo sapiens folate receptor 1 (adult) (FOLR1), transcript variant 2, mRNA
NM_006991	Homo sapiens zinc finger protein 197 (ZNF197), mRNA
NM_018946	Homo sapiens N-acetylneuraminic acid phosphate synthase; sialic acid synthase
_	(SAS), mRNA
NM 003979	Homo sapiens retinoic acid induced 3 (RAI3), mRNA
NM 021785	Homo sapiens retinoic acid induced 2 (RAI2), mRNA
NM 001436	Homo sapiens fibrillarin (FBL), mRNA
NM 012151	Homo sapiens coagulation factor VIII-associated (intronic transcript) (F8A),
1.2	mRNA
NM 007170	Homo sapiens testis-specific kinase 2 (TESK2), mRNA
	1 captono toono opoonio minaso a (12012), matin

NM_006285	II regions to the service 1 (TECK1) DNIA
	Homo sapiens testis-specific kinase 1 (TESK1), mRNA
NM 016424	Homo sapiens cisplatin resistance-associated overexpressed protein (LUC7A),
	mRNA
NM_012152	Homo sapiens endothelial differentiation, lysophosphatidic acid G-protein-
	coupled receptor, 7 (EDG7), mRNA
NM_007360	Homo sapiens DNA segment on chromosome 12 (unique) 2489 expressed
_	sequence (D12S2489E), mRNA
NM_004924	Homo sapiens actinin, alpha 4 (ACTN4), mRNA
NM 001102	Homo sapiens actinin, alpha 1 (ACTN1), mRNA
NM 012128	Homo sapiens chloride channel, calcium activated, family member 4 (CLCA4),
	mRNA
NM 014551	Homo sapiens hypothetical protein 384D8_6 (384D8-2), mRNA
NM_018977	Homo sapiens neuroligin 3 (NLGN3), mRNA
NM 001103	Homo sapiens actinin, alpha 2 (ACTN2), mRNA
NM 022569	Homo sapiens N-deacetylase/N-sulfotransferase 4 (NDST4), mRNA
NM 005892	Homo sapiens formin-like (FMNL), mRNA
NM 016370	Homo sapiens RAB9-like protein (RAB9L), mRNA
NM 012135	Homo sapiens DNA segment on chromosome 6(unique) 2654 expressed
_	sequence (D6S2654E), mRNA
NM 007161	Homo sapiens DNA segment on chromosome 6 (unique) 49 expressed sequence,
	NK cell triggering receptor, p30 (D6S49E), mRNA
NM_006114	Homo sapiens DNA segment on chromosome 19 (unique) 1177 expressed
_	sequence (D19S1177E), mRNA
NM 006014	Homo sapiens DNA segment on chromosome X (unique) 9879 expressed
_	sequence (DXS9879E), mRNA
NM 004699	Homo sapiens DNA segment on chromosome X (unique) 9928 expressed
_	sequence (DXS9928E), mRNA
NM_003683	Homo sapiens DNA segment on chromosome 21 (unique) 2056 expressed
	sequence (D21S2056E), mRNA
NM_015484	Homo sapiens GCIP-interacting protein p29 (P29), mRNA
NM_013263	Homo sapiens bromodomain-containing 7 (BRD7), mRNA
NM_022157	Homo sapiens Rag C protein (GTR2), mRNA
NM_014604	Homo sapiens Tax interaction protein 1 (TIP-1), mRNA
NM_001915	Homo sapiens cytochrome b-561 (CYB561), mRNA
NM_012188	Homo sapiens forkhead box I1 (FOXI1), mRNA
NM_016148	Homo sapiens somatostatin receptor-interacting protein (SSTRIP), mRNA
NM_022482	Homo sapiens hypothetical protein FLJ21794 (FLJ21794), mRNA
NM_022493	Homo sapiens hypothetical protein FLJ21988 (FLJ21988), mRNA
NM_022489	Homo sapiens hypothetical protein FLJ22056 (FLJ22056), mRNA
NM_022485	Homo sapiens hypothetical protein FLJ22405 (FLJ22405), mRNA
NM_022464	Homo sapiens endoplasmic reticulum chaperone SIL1, homolog of yeast (SIL1),
_	mRNA
NM 022456	Homo sapiens hypothetical protein FLJ22548 similar to gene trap PAT 12
	(FLJ22548), mRNA
NM_022450	Homo sapiens hypothetical protein FLJ22357 similar to epidermal growth factor
_	receptor-related protein (FLJ22357), mRNA
NM 022443	Homo sapiens myeloid leukemia factor 1 (MLF1), mRNA
NM 022136	Homo sapiens SAM domain, SH3 domain and nuclear localisation signals, 1
	(SAMSN1), mRNA
NDA 012217	Homo sapiens mast cell tryptase (TPSD1), mRNA
19191 01221/ 1	
NM_012217 NM_020366	Homo sapiens retinitis pigmentosa GTPase regulator interacting protein 1

mRNA NM_004885 Homo sapiens neuropeptide G protein-coupled receptor; neuropeptide FF 2 (NPGPR), mRNA NM_002958 Homo sapiens RYK receptor-like tyrosine kinase (RYK), mRNA NM_002911 Homo sapiens reversion-inducing-cysteine-rich protein with kazal motifs (RECK), mRNA NM_001111 Homo sapiens archain 1 (ARCN1), mRNA NM_001659 Homo sapiens archain 1 (ARCN1), mRNA NM_001669 Homo sapiens scrin-like 7B (ACTL7B), mRNA NM_006686 Homo sapiens actin-like 7A (ACTL7A), mRNA NM_005856 Homo sapiens receptor (calcitonin) activity modifying protein 3 (RAMP3), mRNA NM_005854 Homo sapiens receptor (calcitonin) activity modifying protein 2 (RAMP2), mRNA NM_005855 Homo sapiens receptor (calcitonin) activity modifying protein 1 (RAMP1), mRNA NM_005856 Homo sapiens receptor (calcitonin) activity modifying protein 1 (RAMP1), mRNA NM_005856 Homo sapiens receptor (calcitonin) activity modifying protein 1 (RAMP1), mRNA NM_005856 Homo sapiens receptor (calcitonin) activity modifying protein 1 (RAMP1), mRNA NM_005856 Homo sapiens RAN binding protein 9 (RANBP9), mRNA NM_000473 Homo sapiens RAN binding protein 9 (RANBP9), mRNA NM_0004634 Homo sapiens shormodomain and PHD finger containing, 1 (BRPF1), mRNA NM_000140 Homo sapiens ferrochelatase (protopr)hyria) (FECH), nuclear gene encoding mitochondrial protein, mRNA NM_000027 Homo sapiens aspartylglucosaminidase (AGA), mRNA NM_000026 Homo sapiens adenylosuccinate lyase (ADSL), mRNA NM_000027 Homo sapiens adenylosuccinate lyase (ADSL), mRNA NM_000026 Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA NM_000017 Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA NM_000017 Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA NM_000016 Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA NM_000017 Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA NM_000018 Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA NM_00017 Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA NM_00018 Homo sapiens activin A receptor type II-like 1 (ACVRL1		
NM_014946   Homo sapiens spastic paraplegia 4 (autosomal dominant; spastin) (SPG4), mRNA   mRNA   Homo sapiens neuropeptide FF 1; RFamide-related peptide receptor (OT7T022 mRNA   NM_004885   Homo sapiens neuropeptide G protein-coupled receptor; neuropeptide FF 2 (NPGPR), mRNA   NM_002958   Homo sapiens RYK receptor-like tyrosine kinase (RYK), mRNA   NM_002951   Homo sapiens RYK receptor-like tyrosine kinase (RYK), mRNA   NM_002951   Homo sapiens reversion-inducing-cysteine-rich protein with kazal motifs (RECK), mRNA   Homo sapiens reversion-inducing-cysteine-rich protein with kazal motifs (RECK), mRNA   Homo sapiens strohain 1 (ARCN1), mRNA   NM_016639   Homo sapiens archain 1 (ARCN1), mRNA   NM_006687   Homo sapiens actin-like 7B (ACTL7B), mRNA   NM_005856   Homo sapiens actin-like 7B (ACTL7B), mRNA   NM_005856   Homo sapiens receptor (calcitonin) activity modifying protein 3 (RAMP3), mRNA   NM_005854   Homo sapiens receptor (calcitonin) activity modifying protein 1 (RAMP1), mRNA   NM_005403   Homo sapiens receptor (calcitonin) activity modifying protein 1 (RAMP1), mRNA   NM_004634   Homo sapiens receptor (calcitonin) activity modifying protein 1 (RAMP1), mRNA   NM_004634   Homo sapiens muclear receptor subfamily 0, group B, member 1 (NR0B1), mRNA   NM_004634   Homo sapiens fronchelatase (protoporphyria) (PECH), nuclear gene encoding mitochondrial protein, mRNA   NM_000464   Homo sapiens samnolevulinate, delta-, dehydratase (ALAD), mRNA   NM_000020   Homo sapiens aspimolevulinate, delta-, dehydratase (ALAD), mRNA   NM_000020   Homo sapiens aspimolevulinate, delta-, dehydratase (ALAD), mRNA   NM_000020   Homo sapiens aspimolevulinate, delta-, dehydratase (ALAD), mRNA   NM_000026   Homo sapiens acetyl-Coenzyme A dehydrogenase, C-2 to C-3 short chain (ACADS), nuclear gene encoding mitochondrial protein, mRNA   NM_000026   Homo sapiens acetyl-Coenzyme A dehydrogenase, C-2 to C-3 short chain (ACADS), nuclear gene encoding mitochondrial protein, mRNA   Homo sapiens acetyl-Coenzyme A dehydrogenase, C-2 to C-3 short		
mRNA NM_004885		
mRNA NM_002958 Homo sapiens reversion-inducing-cysteine-rich protein with kazal motifs (RECK), mRNA NM_00291111 Homo sapiens ring finger protein 1 (RNG1), mRNA NM_001111 Homo sapiens archain 1 (ARCN1), mRNA NM_001655 Homo sapiens archain 1 (ARCN1), mRNA NM_001657 Homo sapiens archain 1 (ARCN1), mRNA NM_0016680 Homo sapiens stype 1 transmembrane protein Fn14 (FN14), mRNA NM_006681 Homo sapiens actin-like 7B (ACTL7B), mRNA NM_005856 Homo sapiens actin-like 7A (ACTL7A), mRNA NM_005856 Homo sapiens receptor (calcitonin) activity modifying protein 3 (RAMP3), mRNA NM_005854 Homo sapiens receptor (calcitonin) activity modifying protein 2 (RAMP2), mRNA NM_005855 Homo sapiens receptor (calcitonin) activity modifying protein 1 (RAMP1), mRNA NM_005856 Homo sapiens receptor (calcitonin) activity modifying protein 1 (RAMP1), mRNA NM_005856 Homo sapiens receptor (calcitonin) activity modifying protein 1 (RAMP1), mRNA NM_005856 Homo sapiens receptor (calcitonin) activity modifying protein 1 (RAMP1), mRNA NM_005856 Homo sapiens receptor (calcitonin) activity modifying protein 1 (RAMP1), mRNA NM_000475 Homo sapiens RAN binding protein 9 (RANBP9), mRNA NM_0004634 Homo sapiens RAN binding protein 9 (RANBP9), mRNA NM_0004634 Homo sapiens bromodomain and PHD finger containing, 1 (BRPF1), mRNA NM_000140 Homo sapiens appartylglucosamimidase (AGA), mRNA NM_000021 Homo sapiens adenylosuccinate lyase (ADSL), mRNA NM_000025 Homo sapiens adenylosuccinate lyase (ADSL), mRNA NM_000026 Homo sapiens adenylosuccinate lyase (ADSL), mRNA NM_000017 Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA NM_000018 Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA NM_000016 Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA NM_000017 Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA NM_000018 Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA NM_00017 Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA NM_000018 Homo sapiens erecoding mitochondrial protein, mRNA NM_00018 Homo sapiens act	NM_014946	mRNA
NM_004885 Homo sapiens neuropeptide G protein-coupled receptor; neuropeptide FF 2 (NPGPR), mRNA NM_002931 Homo sapiens RYK receptor-like tyrosine kinase (RYK), mRNA NM_002931 Homo sapiens ring finger protein 1 (RING1), mRNA NM_01111 Homo sapiens reversion-inducing-cysteine-rich protein with kazal motifs (RECK), mRNA NM_016635 Homo sapiens archain 1 (ARCN1), mRNA NM_016636 Homo sapiens stril-like 7B (ACTL/B), mRNA NM_006686 Homo sapiens actin-like 7B (ACTL/B), mRNA NM_006687 Homo sapiens actin-like 7A (ACTL/TA), mRNA NM_005856 Homo sapiens receptor (calcitonin) activity modifying protein 3 (RAMP3), mRNA NM_005854 Homo sapiens receptor (calcitonin) activity modifying protein 2 (RAMP2), mRNA NM_005855 Homo sapiens receptor (calcitonin) activity modifying protein 1 (RAMP1), mRNA NM_004635 Homo sapiens receptor (calcitonin) activity modifying protein 1 (RAMP1), mRNA NM_00463 Homo sapiens receptor subfamily 0, group B, member 1 (NR0B1), mRNA NM_00464 Homo sapiens RAN binding protein 9 (RANBP9), mRNA NM_00464 Homo sapiens serrochelatase (protoporphyria) (FECH), nuclear gene encoding mitochondrial protein, mRNA NM_00001 Homo sapiens sapirylglucosaminidase (AGA), mRNA NM_00002 Homo sapiens adentylosuccinate lyase (ADSL), mRNA NM_00002 Homo sapiens adentylosuccinate lyase (ADSL), mRNA NM_00001 Homo sapiens activin A receptor type E-like 1 (ACVEL1), mRNA NM_00001 Homo sapiens activin A receptor type E-like 1 (ACVEL1), mRNA NM_00001 Homo sapiens activin A receptor type E-like 1 (ACVEL1), mRNA NM_00001 Homo sapiens activin A receptor type E-like 1 (ACVEL1), mRNA NM_00001 Homo sapiens activin A receptor type E-like 1 (ACVEL1), mRNA NM_00001 Homo sapiens activin A receptor type E-like 1 (ACVEL1), mRNA NM_00001 Homo sapiens activin A receptor type E-like 1 (ACVEL1), mRNA NM_00017 Homo sapiens activin A receptor type E-like 1 (ACVEL1), mRNA NM_00018 Homo sapiens activin A receptor type E-like 1 (ACVEL1), mRNA NM_00018 Homo sapiens activin A receptor type E-like 1 (ACVEL1), mRNA NM_00018 Homo sapiens spiral decompliant to mouse	NM_022146	Homo sapiens neuropeptide FF 1; RFamide-related peptide receptor (OT7T022), mRNA
NM	NM_004885	Homo sapiens neuropeptide G protein-coupled receptor; neuropeptide FF 2
NM   002111	NM 002958	
NM_021111		Homo sapiens ring finger protein 1 (RING1), mRNA
NM		Homo sapiens reversion-inducing-cysteine-rich protein with kazal motifs
NM         006686         Homo sapiens actin-like 7B (ACTL7B), mRNA           NM         006687         Homo sapiens actin-like 7A (ACTL7A), mRNA           NM_005856         Homo sapiens receptor (calcitonin) activity modifying protein 3 (RAMP3), mRNA           NM_005854         Homo sapiens receptor (calcitonin) activity modifying protein 2 (RAMP2), mRNA           NM_005855         Homo sapiens receptor (calcitonin) activity modifying protein 1 (RAMP1), mRNA           NM_000475         Homo sapiens nuclear receptor subfamily 0, group B, member 1 (NR0B1), mRNA           NM_005493         Homo sapiens RAN binding protein 9 (RANBP9), mRNA           NM_004634         Homo sapiens bromodomain and PHD finger containing, 1 (BRPF1), mRNA           NM_000140         Homo sapiens ferrochelatase (protoporphyria) (FECH), nuclear gene encoding mitochondrial protein, mRNA           NM_000021         Homo sapiens aminolevulinate, delta-, dehydratase (ALAD), mRNA           NM_000025         Homo sapiens adrenergic, beta-3-, receptor (ADRB3), mRNA           NM_000026         Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA           NM_000019         Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA           NM_000010         Homo sapiens acyl-Coenzyme A dehydrogenase, very long chain (ACADVL) nuclear gene encoding mitochondrial protein, mRNA           NM_00018         Homo sapiens acyl-Coenzyme A dehydrogenase, C-2 to C-3 short chain (ACADM), nuclear gene encoding mitoch	NM_001655	Homo sapiens archain 1 (ARCN1), mRNA
NM 00687   Homo sapiens actin-like 7A (ACTL7A), mRNA   NM 005856   Homo sapiens receptor (calcitonin) activity modifying protein 3 (RAMP3), mRNA   NM 005855   Homo sapiens receptor (calcitonin) activity modifying protein 2 (RAMP2), mRNA   NM 005855   Homo sapiens receptor (calcitonin) activity modifying protein 1 (RAMP1), mRNA   NM 000475   Homo sapiens nuclear receptor subfamily 0, group B, member 1 (NR0B1), mRNA   NM 005493   Homo sapiens RAN binding protein 9 (RANBP9), mRNA   NM 004634   Homo sapiens bromodomain and PHD finger containing, 1 (BRPF1), mRNA   NM 000140   Homo sapiens ferrochelatase (protoporphyria) (FECH), nuclear gene encoding mitochondrial protein, mRNA   NM 000021   Homo sapiens aminolevulinate, delta-, dehydratase (ALAD), mRNA   NM 000025   Homo sapiens adenylosuccinate lyase (ADSL), mRNA   NM 000026   Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA   NM 000020   Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA   NM 000018   Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA   NM 000016   Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA   NM 000016   Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA   NM 000016   Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA   NM 000016   Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA   NM 000016   Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA   NM 000016   Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA   NM 000016   Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA   NM 000016   Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA   NM 000016   Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA   NM 000016   Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA   NM 000016   Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA   NM 000016   Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA   Homo sapiens activition type II-like 1 (ACVRL1), mRNA   NM 000016   Homo sapiens	NM 016639	Homo sapiens type I transmembrane protein Fn14 (FN14), mRNA
NM         006687         Homo sapiens actin-like 7A (ACTL7A), mRNA           NM_005856         Homo sapiens receptor (calcitonin) activity modifying protein 3 (RAMP3), mRNA           NM_005854         Homo sapiens receptor (calcitonin) activity modifying protein 2 (RAMP2), mRNA           NM_005855         Homo sapiens receptor (calcitonin) activity modifying protein 1 (RAMP1), mRNA           NM_000475         Homo sapiens nuclear receptor subfamily 0, group B, member 1 (NR0B1), mRNA           NM_005493         Homo sapiens RAN binding protein 9 (RANBP9), mRNA           NM_000440         Homo sapiens bromodomain and PHD finger containing, 1 (BRPF1), mRNA           NM_000140         Homo sapiens afterrochelatase (protoporphyria) (FECH), nuclear gene encoding mitochondrial protein, mRNA           NM_000021         Homo sapiens aminolevulinate, delta-, dehydratase (ALAD), mRNA           NM_000022         Homo sapiens adenylosuccinate lyase (ADSL), mRNA           NM_000025         Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA           NM_000019         Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA           NM_000010         Homo sapiens actyl-Coenzyme A dehydrogenase, very long chain (ACADVL) nuclear gene encoding mitochondrial protein, mRNA           NM_000017         Homo sapiens acyl-Coenzyme A dehydrogenase, C-2 to C-3 short chain (ACADM), nuclear gene encoding mitochondrial protein, mRNA           NM_000016         Homo sapiens adenylate kinase 1 (AK1), mR	NM 006686	Homo sapiens actin-like 7B (ACTL7B), mRNA
NM_005856 Homo sapiens receptor (calcitonin) activity modifying protein 3 (RAMP3), mRNA  NM_005854 Homo sapiens receptor (calcitonin) activity modifying protein 2 (RAMP2), mRNA  NM_005855 Homo sapiens receptor (calcitonin) activity modifying protein 1 (RAMP1), mRNA  NM_000475 Homo sapiens nuclear receptor subfamily 0, group B, member 1 (NR0B1), mRNA  NM_005493 Homo sapiens RAN binding protein 9 (RANBP9), mRNA  NM_004634 Homo sapiens bromodomain and PHD finger containing, 1 (BRPF1), mRNA  NM_000140 Homo sapiens ferrochelatase (protoporphyria) (FECH), nuclear gene encoding mitochondrial protein, mRNA  NM_000027 Homo sapiens aminolevulinate, delta-, dehydratase (ALAD), mRNA  NM_000026 Homo sapiens adenylosuccinate lyase (ADSL), mRNA  NM_000025 Homo sapiens adrenergic, beta-3-, receptor (ADRB3), mRNA  NM_000020 Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA  NM_000019 Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA  NM_000018 Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA  NM_000018 Homo sapiens acyl-Coenzyme A dehydrogenase, very long chain (ACADVL) nuclear gene encoding mitochondrial protein, mRNA  NM_000016 Homo sapiens acyl-Coenzyme A dehydrogenase, c-2 to C-3 short chain (ACADS), nuclear gene encoding mitochondrial protein, mRNA  NM_000016 Homo sapiens acyl-Coenzyme A dehydrogenase, C-2 to C-3 short chain (ACADM), nuclear gene encoding mitochondrial protein, mRNA  NM_00016 Homo sapiens acyl-Coenzyme A dehydrogenase, C-2 to C-3 short chain (ACADM), nuclear gene encoding mitochondrial protein, mRNA  NM_00030 Homo sapiens acyl-Coenzyme A dehydrogenase, C-2 to C-3 short chain (ACADM), nuclear gene encoding mitochondrial protein, mRNA  NM_0030 Homo sapiens acyl-Coenzyme A dehydrogenase, C-2 to C-3 short chain (ACADM), nuclear gene encoding mitochondrial protein, mRNA  NM_0030 Homo sapiens acyl-Coenzyme A dehydrogenase, C-2 to C-3 short chain (ACADM), nuclear gene encoding mitochondrial protein, mRNA  NM_00300 Homo sapiens acyl-Coenzyme A dehydrogenase, C-2 to C-12 straigh		
mRNA NM_005855 Homo sapiens receptor (calcitonin) activity modifying protein 1 (RAMP1), mRNA NM_000475 Homo sapiens nuclear receptor subfamily 0, group B, member 1 (NR0B1), mRNA NM_005493 Homo sapiens RAN binding protein 9 (RANBP9), mRNA NM_004634 Homo sapiens bromodomain and PHD finger containing, 1 (BRPF1), mRNA NM_000140 Homo sapiens ferrochelatase (protoporphyria) (FECH), nuclear gene encoding mitochondrial protein, mRNA NM_000021 Homo sapiens aminolevulinate, delta-, dehydratase (ALAD), mRNA NM_000027 Homo sapiens aspartylglucosaminidase (AGA), mRNA NM_000026 Homo sapiens adenylosuccinate lyase (ADSL), mRNA NM_000026 Homo sapiens aderenergic, beta-3-, receptor (ADRB3), mRNA NM_000020 Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA NM_000019 Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA NM_000018 Homo sapiens acyl-Coenzyme A acetyltransferase 1 (acetoacetyl Coenzyme thiolase) (ACAT1), nuclear gene encoding mitochondrial protein, mRNA NM_000017 Homo sapiens acyl-Coenzyme A dehydrogenase, very long chain (ACADVL) nuclear gene encoding mitochondrial protein, mRNA NM_000016 Homo sapiens acyl-Coenzyme A dehydrogenase, C-2 to C-3 short chain (ACADM), nuclear gene encoding mitochondrial protein, mRNA NM_000016 Homo sapiens acyl-Coenzyme A dehydrogenase, C-2 to C-12 straight chain (ACADM), nuclear gene encoding mitochondrial protein, mRNA NM_000476 Homo sapiens adenylate kinase 1 (AK1), mRNA NM_001830 Homo sapiens shypothetical protein similar to mouse Dnajl1 (DNAJL1), mRNA NM_022305 Homo sapiens aminopeptidase (LOC64167), mRNA NM_022305 Homo sapiens aminopeptidase (LOC64167), mRNA NM_022306 Homo sapiens growth differentiation factor 8 (GDF8), mRNA NM_001789 Homo sapiens FXYD domain-containing ion transport regulator 7 (FXYD7),		Homo sapiens receptor (calcitonin) activity modifying protein 3 (RAMP3),
mRNA NM_000475 Homo sapiens nuclear receptor subfamily 0, group B, member 1 (NR0B1), mRNA NM_005493 Homo sapiens RAN binding protein 9 (RANBP9), mRNA NM_004634 Homo sapiens bromodomain and PHD finger containing, 1 (BRPF1), mRNA NM_000140 Homo sapiens ferrochelatase (protoporphyria) (FECH), nuclear gene encoding mitochondrial protein, mRNA NM_000021 Homo sapiens aminolevulinate, delta-, dehydratase (ALAD), mRNA NM_000025 Homo sapiens adenylosuccinate lyase (ADSL), mRNA NM_000026 Homo sapiens adrenergic, beta-3-, receptor (ADRB3), mRNA NM_000020 Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA NM_000019 Homo sapiens acetyl-Coenzyme A acetyltransferase 1 (acetoacetyl Coenzyme thiolase) (ACAT1), nuclear gene encoding mitochondrial protein, mRNA NM_000018 Homo sapiens acyl-Coenzyme A dehydrogenase, very long chain (ACADVL) nuclear gene encoding mitochondrial protein, mRNA NM_000017 Homo sapiens acyl-Coenzyme A dehydrogenase, C-2 to C-3 short chain (ACADS), nuclear gene encoding mitochondrial protein, mRNA NM_000016 Homo sapiens acyl-Coenzyme A dehydrogenase, C-2 to C-3 short chain (ACADM), nuclear gene encoding mitochondrial protein, mRNA NM_000016 Homo sapiens acyl-Coenzyme A dehydrogenase, C-4 to C-12 straight chain (ACADM), nuclear gene encoding mitochondrial protein, mRNA NM_00016 Homo sapiens adenylate kinase 1 (AK1), mRNA NM_001830 Homo sapiens adenylate kinase 1 (AK1), mRNA NM_002335 Homo sapiens shypothetical protein similar to mouse Dnajl1 (DNAJL1), mRNA NM_02235 Homo sapiens aminopeptidase (LOC64167), mRNA NM_002335 Homo sapiens sminopeptidase (LOC64167), mRNA NM_001789 Homo sapiens cell division cycle 25A (CDC25A), mRNA NM_0022006 Homo sapiens FXYD domain-containing ion transport regulator 7 (FXYD7),	NM_005854	
mRNA  NM 005493 Homo sapiens RAN binding protein 9 (RANBP9), mRNA  NM 004634 Homo sapiens bromodomain and PHD finger containing, 1 (BRPF1), mRNA  NM 000140 Homo sapiens ferrochelatase (protoporphyria) (FECH), nuclear gene encoding mitochondrial protein, mRNA  NM 000031 Homo sapiens aspartylglucosaminidase (AGA), mRNA  NM 000027 Homo sapiens adenylosuccinate lyase (ADSL), mRNA  NM 000026 Homo sapiens adrenergic, beta-3-, receptor (ADRB3), mRNA  NM 000020 Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA  NM 000010 Homo sapiens activl-Coenzyme A acetyltransferase 1 (acetoacetyl Coenzyme thiolase) (ACAT1), nuclear gene encoding mitochondrial protein, mRNA  NM 000018 Homo sapiens acyl-Coenzyme A dehydrogenase, very long chain (ACADVL) nuclear gene encoding mitochondrial protein, mRNA  NM 000017 Homo sapiens acyl-Coenzyme A dehydrogenase, C-2 to C-3 short chain (ACADM), nuclear gene encoding mitochondrial protein, mRNA  NM 000016 Homo sapiens acyl-Coenzyme A dehydrogenase, C-4 to C-12 straight chain (ACADM), nuclear gene encoding mitochondrial protein, mRNA  NM 000016 Homo sapiens acyl-Coenzyme A dehydrogenase, C-4 to C-12 straight chain (ACADM), nuclear gene encoding mitochondrial protein, mRNA  NM 00016 Homo sapiens acyl-Coenzyme A dehydrogenase, C-4 to C-12 straight chain (ACADM), nuclear gene encoding mitochondrial protein, mRNA  NM 00017 Homo sapiens acyl-Coenzyme A dehydrogenase, C-4 to C-12 straight chain (ACADM), nuclear gene encoding mitochondrial protein, mRNA  NM 000176 Homo sapiens hypothetical protein similar to mouse Dnajl1 (DNAJL1), mRN  NM 001830 Homo sapiens bypothetical protein similar to mouse Dnajl1 (DNAJL1), mRN  NM 022335 Homo sapiens spyothetical protein PRO2849 (PRO2849), mRNA  NM 001789 Homo sapiens cell division cycle 25A (CDC25A), mRNA  NM 001789 Homo sapiens FXYD domain-containing ion transport regulator 7 (FXYD7),	NM_005855	
NM_004634Homo sapiens bromodomain and PHD finger containing, 1 (BRPF1), mRNANM_000140Homo sapiens ferrochelatase (protoporphyria) (FECH), nuclear gene encoding mitochondrial protein, mRNANM_000031Homo sapiens aminolevulinate, delta-, dehydratase (ALAD), mRNANM_000027Homo sapiens aspartylglucosaminidase (AGA), mRNANM_000026Homo sapiens adenylosuccinate lyase (ADSL), mRNANM_000025Homo sapiens adrenergic, beta-3-, receptor (ADRB3), mRNANM_000020Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNANM_000019Homo sapiens actyl-Coenzyme A acetyltransferase 1 (acetoacetyl Coenzyme thiolase) (ACAT1), nuclear gene encoding mitochondrial protein, mRNANM_000018Homo sapiens acyl-Coenzyme A dehydrogenase, very long chain (ACADVL) nuclear gene encoding mitochondrial protein, mRNANM_000017Homo sapiens acyl-Coenzyme A dehydrogenase, C-2 to C-3 short chain (ACADS), nuclear gene encoding mitochondrial protein, mRNANM_000016Homo sapiens acyl-Coenzyme A dehydrogenase, C-4 to C-12 straight chain (ACADM), nuclear gene encoding mitochondrial protein, mRNANM_000476Homo sapiens adenylate kinase 1 (AK1), mRNANM_001830Homo sapiens hypothetical protein similar to mouse Dnajl1 (DNAJL1), mRN, NM, 022365Homo sapiens hypothetical protein PRO2849 (PRO2849), mRNANM_022350Homo sapiens shypothetical protein PRO2849 (PRO2849), mRNANM_001789Homo sapiens scell division cycle 25A (CDC25A), mRNANM_001789Homo sapiens FXYD domain-containing ion transport regulator 7 (FXYD7),	NM_000475	
NM_004634Homo sapiens bromodomain and PHD finger containing, 1 (BRPF1), mRNANM_000140Homo sapiens ferrochelatase (protoporphyria) (FECH), nuclear gene encoding mitochondrial protein, mRNANM_000031Homo sapiens aminolevulinate, delta-, dehydratase (ALAD), mRNANM_000027Homo sapiens aspartylglucosaminidase (AGA), mRNANM_000026Homo sapiens adenylosuccinate lyase (ADSL), mRNANM_000025Homo sapiens adrenergic, beta-3-, receptor (ADRB3), mRNANM_000020Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNANM_000019Homo sapiens actyl-Coenzyme A acetyltransferase 1 (acetoacetyl Coenzyme thiolase) (ACAT1), nuclear gene encoding mitochondrial protein, mRNANM_000018Homo sapiens acyl-Coenzyme A dehydrogenase, very long chain (ACADVL) nuclear gene encoding mitochondrial protein, mRNANM_000017Homo sapiens acyl-Coenzyme A dehydrogenase, C-2 to C-3 short chain (ACADS), nuclear gene encoding mitochondrial protein, mRNANM_000016Homo sapiens acyl-Coenzyme A dehydrogenase, C-4 to C-12 straight chain (ACADM), nuclear gene encoding mitochondrial protein, mRNANM_001830Homo sapiens adenylate kinase 1 (AK1), mRNANM_002365Homo sapiens hypothetical protein similar to mouse Dnajl1 (DNAJL1), mRN, NM_022350Homo sapiens hypothetical protein PRO2849 (PRO2849), mRNANM_005259Homo sapiens growth differentiation factor 8 (GDF8), mRNANM_001789Homo sapiens FXYD domain-containing ion transport regulator 7 (FXYD7),	NM 005493	Homo sapiens RAN binding protein 9 (RANBP9), mRNA
mitochondrial protein, mRNA  NM_000031 Homo sapiens aminolevulinate, delta-, dehydratase (ALAD), mRNA  NM_000027 Homo sapiens aspartylglucosaminidase (AGA), mRNA  NM_000026 Homo sapiens adenylosuccinate lyase (ADSL), mRNA  NM_000025 Homo sapiens adrenergic, beta-3-, receptor (ADRB3), mRNA  NM_000020 Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA  NM_000019 Homo sapiens acetyl-Coenzyme A acetyltransferase 1 (acetoacetyl Coenzyme thiolase) (ACAT1), nuclear gene encoding mitochondrial protein, mRNA  NM_000018 Homo sapiens acyl-Coenzyme A dehydrogenase, very long chain (ACADVL) nuclear gene encoding mitochondrial protein, mRNA  NM_000017 Homo sapiens acyl-Coenzyme A dehydrogenase, C-2 to C-3 short chain (ACADS), nuclear gene encoding mitochondrial protein, mRNA  NM_000016 Homo sapiens acyl-Coenzyme A dehydrogenase, C-4 to C-12 straight chain (ACADM), nuclear gene encoding mitochondrial protein, mRNA  NM_000476 Homo sapiens adenylate kinase 1 (AK1), mRNA  NM_001830 Homo sapiens chloride channel 4 (CLCN4), mRNA  NM_02365 Homo sapiens hypothetical protein similar to mouse Dnajl1 (DNAJL1), mRN, NM_02350 Homo sapiens aminopeptidase (LOC64167), mRNA  NM_022350 Homo sapiens provth differentiation factor 8 (GDF8), mRNA  NM_001789 Homo sapiens cell division cycle 25A (CDC25A), mRNA  NM_002006 Homo sapiens FXYD domain-containing ion transport regulator 7 (FXYD7),	NM 004634	Homo sapiens bromodomain and PHD finger containing, 1 (BRPF1), mRNA
NM_000031 Homo sapiens aminolevulinate, delta-, dehydratase (ALAD), mRNA NM_000027 Homo sapiens aspartylglucosaminidase (AGA), mRNA NM_000026 Homo sapiens adenylosuccinate lyase (ADSL), mRNA NM_000025 Homo sapiens adrenergic, beta-3-, receptor (ADRB3), mRNA NM_000020 Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA NM_000019 Homo sapiens acetyl-Coenzyme A acetyltransferase 1 (acetoacetyl Coenzyme thiolase) (ACAT1), nuclear gene encoding mitochondrial protein, mRNA NM_000018 Homo sapiens acyl-Coenzyme A dehydrogenase, very long chain (ACADVL) nuclear gene encoding mitochondrial protein, mRNA NM_000017 Homo sapiens acyl-Coenzyme A dehydrogenase, C-2 to C-3 short chain (ACADS), nuclear gene encoding mitochondrial protein, mRNA NM_000016 Homo sapiens acyl-Coenzyme A dehydrogenase, C-4 to C-12 straight chain (ACADM), nuclear gene encoding mitochondrial protein, mRNA NM_000476 Homo sapiens adenylate kinase 1 (AK1), mRNA NM_001830 Homo sapiens chloride channel 4 (CLCN4), mRNA NM_02365 Homo sapiens hypothetical protein similar to mouse Dnajl1 (DNAJL1), mRN, NM_02350 Homo sapiens aminopeptidase (LOC64167), mRNA NM_02335 Homo sapiens provth differentiation factor 8 (GDF8), mRNA NM_001789 Homo sapiens cell division cycle 25A (CDC25A), mRNA NM_002006 Homo sapiens FXYD domain-containing ion transport regulator 7 (FXYD7),	NM_000140	Homo sapiens ferrochelatase (protoporphyria) (FECH), nuclear gene encoding mitochondrial protein, mRNA
NM_000027 Homo sapiens aspartylglucosaminidase (AGA), mRNA NM_000026 Homo sapiens adenylosuccinate lyase (ADSL), mRNA NM_000025 Homo sapiens adrenergic, beta-3-, receptor (ADRB3), mRNA NM_000020 Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA NM_000019 Homo sapiens acetyl-Coenzyme A acetyltransferase 1 (acetoacetyl Coenzyme thiolase) (ACAT1), nuclear gene encoding mitochondrial protein, mRNA NM_000018 Homo sapiens acyl-Coenzyme A dehydrogenase, very long chain (ACADVL) nuclear gene encoding mitochondrial protein, mRNA NM_000017 Homo sapiens acyl-Coenzyme A dehydrogenase, C-2 to C-3 short chain (ACADS), nuclear gene encoding mitochondrial protein, mRNA NM_000016 Homo sapiens acyl-Coenzyme A dehydrogenase, C-4 to C-12 straight chain (ACADM), nuclear gene encoding mitochondrial protein, mRNA NM_000476 Homo sapiens adenylate kinase 1 (AK1), mRNA NM_001830 Homo sapiens chloride channel 4 (CLCN4), mRNA NM_022365 Homo sapiens hypothetical protein similar to mouse Dnajl1 (DNAJL1), mRN, NM_022350 Homo sapiens aminopeptidase (LOC64167), mRNA NM_022351 Homo sapiens growth differentiation factor 8 (GDF8), mRNA NM_001789 Homo sapiens FXYD domain-containing ion transport regulator 7 (FXYD7), NM_022006 Homo sapiens FXYD domain-containing ion transport regulator 7 (FXYD7),	NM 000031	<del></del>
NM_000026 Homo sapiens adenylosuccinate lyase (ADSL), mRNA NM_000025 Homo sapiens adrenergic, beta-3-, receptor (ADRB3), mRNA NM_000020 Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA NM_000019 Homo sapiens acetyl-Coenzyme A acetyltransferase 1 (acetoacetyl Coenzyme thiolase) (ACAT1), nuclear gene encoding mitochondrial protein, mRNA NM_000018 Homo sapiens acyl-Coenzyme A dehydrogenase, very long chain (ACADVL) nuclear gene encoding mitochondrial protein, mRNA NM_000017 Homo sapiens acyl-Coenzyme A dehydrogenase, C-2 to C-3 short chain (ACADS), nuclear gene encoding mitochondrial protein, mRNA NM_000016 Homo sapiens acyl-Coenzyme A dehydrogenase, C-4 to C-12 straight chain (ACADM), nuclear gene encoding mitochondrial protein, mRNA NM_000476 Homo sapiens adenylate kinase 1 (AK1), mRNA NM_001830 Homo sapiens chloride channel 4 (CLCN4), mRNA NM_022365 Homo sapiens hypothetical protein similar to mouse Dnajl1 (DNAJL1), mRN. NM_022350 Homo sapiens aminopeptidase (LOC64167), mRNA NM_022351 Homo sapiens protein prot		
NM_000025Homo sapiens adrenergic, beta-3-, receptor (ADRB3), mRNANM_000020Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNANM_000019Homo sapiens acetyl-Coenzyme A acetyltransferase 1 (acetoacetyl Coenzyme thiolase) (ACAT1), nuclear gene encoding mitochondrial protein, mRNANM_000018Homo sapiens acyl-Coenzyme A dehydrogenase, very long chain (ACADVL) nuclear gene encoding mitochondrial protein, mRNANM_000017Homo sapiens acyl-Coenzyme A dehydrogenase, C-2 to C-3 short chain (ACADS), nuclear gene encoding mitochondrial protein, mRNANM_000016Homo sapiens acyl-Coenzyme A dehydrogenase, C-4 to C-12 straight chain (ACADM), nuclear gene encoding mitochondrial protein, mRNANM_000476Homo sapiens adenylate kinase 1 (AK1), mRNANM_001830Homo sapiens chloride channel 4 (CLCN4), mRNANM_022365Homo sapiens hypothetical protein similar to mouse Dnajl1 (DNAJL1), mRN, NM_022350Homo sapiens aminopeptidase (LOC64167), mRNANM_02235Homo sapiens prowth differentiation factor 8 (GDF8), mRNANM_001789Homo sapiens cell division cycle 25A (CDC25A), mRNANM_02206Homo sapiens FXYD domain-containing ion transport regulator 7 (FXYD7),	NM 000026	<u>, , , , , , , , , , , , , , , , , , , </u>
NM_000020Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNANM_000019Homo sapiens acetyl-Coenzyme A acetyltransferase 1 (acetoacetyl Coenzyme thiolase) (ACAT1), nuclear gene encoding mitochondrial protein, mRNANM_000018Homo sapiens acyl-Coenzyme A dehydrogenase, very long chain (ACADVL) nuclear gene encoding mitochondrial protein, mRNANM_000017Homo sapiens acyl-Coenzyme A dehydrogenase, C-2 to C-3 short chain (ACADS), nuclear gene encoding mitochondrial protein, mRNANM_00016Homo sapiens acyl-Coenzyme A dehydrogenase, C-4 to C-12 straight chain (ACADM), nuclear gene encoding mitochondrial protein, mRNANM_000476Homo sapiens adenylate kinase 1 (AK1), mRNANM_001830Homo sapiens chloride channel 4 (CLCN4), mRNANM_022365Homo sapiens hypothetical protein similar to mouse Dnajl1 (DNAJL1), mRN, nM_022350NM_022350Homo sapiens aminopeptidase (LOC64167), mRNANM_022351Homo sapiens protein PRO2849 (PRO2849), mRNANM_005259Homo sapiens growth differentiation factor 8 (GDF8), mRNANM_001789Homo sapiens Cell division cycle 25A (CDC25A), mRNANM_022006Homo sapiens FXYD domain-containing ion transport regulator 7 (FXYD7),	NM 000025	
NM_000019 Homo sapiens acetyl-Coenzyme A acetyltransferase 1 (acetoacetyl Coenzyme thiolase) (ACAT1), nuclear gene encoding mitochondrial protein, mRNA  NM_000018 Homo sapiens acyl-Coenzyme A dehydrogenase, very long chain (ACADVL) nuclear gene encoding mitochondrial protein, mRNA  NM_000017 Homo sapiens acyl-Coenzyme A dehydrogenase, C-2 to C-3 short chain (ACADS), nuclear gene encoding mitochondrial protein, mRNA  NM_000016 Homo sapiens acyl-Coenzyme A dehydrogenase, C-4 to C-12 straight chain (ACADM), nuclear gene encoding mitochondrial protein, mRNA  NM_000476 Homo sapiens adenylate kinase 1 (AK1), mRNA  NM_001830 Homo sapiens chloride channel 4 (CLCN4), mRNA  NM_022365 Homo sapiens hypothetical protein similar to mouse Dnajl1 (DNAJL1), mRN, NM_022350 Homo sapiens aminopeptidase (LOC64167), mRNA  NM_022335 Homo sapiens hypothetical protein PRO2849 (PRO2849), mRNA  NM_005259 Homo sapiens growth differentiation factor 8 (GDF8), mRNA  NM_001789 Homo sapiens cell division cycle 25A (CDC25A), mRNA  NM_022006 Homo sapiens FXYD domain-containing ion transport regulator 7 (FXYD7),		
NM_000018 Homo sapiens acyl-Coenzyme A dehydrogenase, very long chain (ACADVL) nuclear gene encoding mitochondrial protein, mRNA  NM_000017 Homo sapiens acyl-Coenzyme A dehydrogenase, C-2 to C-3 short chain (ACADS), nuclear gene encoding mitochondrial protein, mRNA  NM_000016 Homo sapiens acyl-Coenzyme A dehydrogenase, C-4 to C-12 straight chain (ACADM), nuclear gene encoding mitochondrial protein, mRNA  NM_000476 Homo sapiens adenylate kinase 1 (AK1), mRNA  NM_01830 Homo sapiens chloride channel 4 (CLCN4), mRNA  NM_022365 Homo sapiens hypothetical protein similar to mouse Dnajl1 (DNAJL1), mRNA  NM_022350 Homo sapiens aminopeptidase (LOC64167), mRNA  NM_022351 Homo sapiens hypothetical protein PRO2849 (PRO2849), mRNA  NM_005259 Homo sapiens growth differentiation factor 8 (GDF8), mRNA  NM_01789 Homo sapiens cell division cycle 25A (CDC25A), mRNA  NM_022006 Homo sapiens FXYD domain-containing ion transport regulator 7 (FXYD7),		Homo sapiens acetyl-Coenzyme A acetyltransferase 1 (acetoacetyl Coenzyme A
NM_000017 Homo sapiens acyl-Coenzyme A dehydrogenase, C-2 to C-3 short chain (ACADS), nuclear gene encoding mitochondrial protein, mRNA  NM_000016 Homo sapiens acyl-Coenzyme A dehydrogenase, C-4 to C-12 straight chain (ACADM), nuclear gene encoding mitochondrial protein, mRNA  NM_000476 Homo sapiens adenylate kinase 1 (AK1), mRNA  NM_001830 Homo sapiens chloride channel 4 (CLCN4), mRNA  NM_022365 Homo sapiens hypothetical protein similar to mouse Dnajl1 (DNAJL1), mRN.  NM_022350 Homo sapiens aminopeptidase (LOC64167), mRNA  NM_022335 Homo sapiens hypothetical protein PRO2849 (PRO2849), mRNA  NM_005259 Homo sapiens growth differentiation factor 8 (GDF8), mRNA  NM_001789 Homo sapiens cell division cycle 25A (CDC25A), mRNA  NM_022006 Homo sapiens FXYD domain-containing ion transport regulator 7 (FXYD7),	NM_000018	Homo sapiens acyl-Coenzyme A dehydrogenase, very long chain (ACADVL),
NM_000016 Homo sapiens acyl-Coenzyme A dehydrogenase, C-4 to C-12 straight chain (ACADM), nuclear gene encoding mitochondrial protein, mRNA  NM_000476 Homo sapiens adenylate kinase 1 (AK1), mRNA  NM_01830 Homo sapiens chloride channel 4 (CLCN4), mRNA  NM_022365 Homo sapiens hypothetical protein similar to mouse Dnajl1 (DNAJL1), mRNA  NM_022350 Homo sapiens aminopeptidase (LOC64167), mRNA  NM_022335 Homo sapiens hypothetical protein PRO2849 (PRO2849), mRNA  NM_005259 Homo sapiens growth differentiation factor 8 (GDF8), mRNA  NM_001789 Homo sapiens cell division cycle 25A (CDC25A), mRNA  NM_022006 Homo sapiens FXYD domain-containing ion transport regulator 7 (FXYD7),	NM_000017	Homo sapiens acyl-Coenzyme A dehydrogenase, C-2 to C-3 short chain
NM_000476 Homo sapiens adenylate kinase 1 (AK1), mRNA  NM_001830 Homo sapiens chloride channel 4 (CLCN4), mRNA  NM_022365 Homo sapiens hypothetical protein similar to mouse Dnajl1 (DNAJL1), mRN.  NM_022350 Homo sapiens aminopeptidase (LOC64167), mRNA  NM_022335 Homo sapiens hypothetical protein PRO2849 (PRO2849), mRNA  NM_005259 Homo sapiens growth differentiation factor 8 (GDF8), mRNA  NM_001789 Homo sapiens cell division cycle 25A (CDC25A), mRNA  NM_022006 Homo sapiens FXYD domain-containing ion transport regulator 7 (FXYD7),	NM_000016	Homo sapiens acyl-Coenzyme A dehydrogenase, C-4 to C-12 straight chain
NM 001830 Homo sapiens chloride channel 4 (CLCN4), mRNA  NM 022365 Homo sapiens hypothetical protein similar to mouse Dnajl1 (DNAJL1), mRN  NM 022350 Homo sapiens aminopeptidase (LOC64167), mRNA  NM 022335 Homo sapiens hypothetical protein PRO2849 (PRO2849), mRNA  NM 005259 Homo sapiens growth differentiation factor 8 (GDF8), mRNA  NM 001789 Homo sapiens cell division cycle 25A (CDC25A), mRNA  NM 022006 Homo sapiens FXYD domain-containing ion transport regulator 7 (FXYD7),	NM 000476	Homo sapiens adenylate kinase 1 (AK1), mRNA
NM_022365 Homo sapiens hypothetical protein similar to mouse Dnajl1 (DNAJL1), mRN. NM_022350 Homo sapiens aminopeptidase (LOC64167), mRNA NM_022335 Homo sapiens hypothetical protein PRO2849 (PRO2849), mRNA NM_005259 Homo sapiens growth differentiation factor 8 (GDF8), mRNA NM_001789 Homo sapiens cell division cycle 25A (CDC25A), mRNA NM_022006 Homo sapiens FXYD domain-containing ion transport regulator 7 (FXYD7),		
NM_022350Homo sapiens aminopeptidase (LOC64167), mRNANM_022335Homo sapiens hypothetical protein PRO2849 (PRO2849), mRNANM_005259Homo sapiens growth differentiation factor 8 (GDF8), mRNANM_001789Homo sapiens cell division cycle 25A (CDC25A), mRNANM_022006Homo sapiens FXYD domain-containing ion transport regulator 7 (FXYD7),		
NM 022335 Homo sapiens hypothetical protein PRO2849 (PRO2849), mRNA NM 005259 Homo sapiens growth differentiation factor 8 (GDF8), mRNA NM 001789 Homo sapiens cell division cycle 25A (CDC25A), mRNA NM 022006 Homo sapiens FXYD domain-containing ion transport regulator 7 (FXYD7),		
NM 005259 Homo sapiens growth differentiation factor 8 (GDF8), mRNA NM 001789 Homo sapiens cell division cycle 25A (CDC25A), mRNA NM 022006 Homo sapiens FXYD domain-containing ion transport regulator 7 (FXYD7),		
NM_001789 Homo sapiens cell division cycle 25A (CDC25A), mRNA NM_022006 Homo sapiens FXYD domain-containing ion transport regulator 7 (FXYD7),		· <del>                                    </del>
NM_022006 Homo sapiens FXYD domain-containing ion transport regulator 7 (FXYD7),		·   · · · · · · · · · · · · · · · · · ·
		mRNA
NM_022003 Homo sapiens FXYD domain-containing ion transport regulator 6 (FXYD6), mRNA	19191_022003	•

NM_020655	Homo sapiens junctophilin 3 (JPH3), mRNA
NM_002855	Homo sapiens poliovirus receptor-related 1 (herpesvirus entry mediator C;
	nectin) (PVRL1), mRNA
NM_012340	Homo sapiens nuclear factor of activated T-cells, cytoplasmic, calcineurin-
	dependent 2 (NFATC2), mRNA
NM_006599	Homo sapiens nuclear factor of activated T-cells 5, tonicity-resonsive (NFAT5),
225	mRNA
NM_006162	Homo sapiens nuclear factor of activated T-cells, cytoplasmic, calcineurin-
NR 6 000061	dependent 1 (NFATC1), mRNA
NM_022061	Homo sapiens ribosomal protein L17 isolog (LOC63875), mRNA
NM_022095	Homo sapiens hypothetical C2H2 zinc finger protein FLJ22504 (FLJ22504),
) Tr 6 000001	mRNA
NM_022091	Homo sapiens dJ467N11.1 protein (DJ467N11.1), mRNA
NM_022084	Homo sapiens hypothetical protein dJ102H19.4 (DJ102H19.4), mRNA
NM_022077	Homo sapiens hypothetical protein dJ1141E15.2 (DJ1141E15.2), mRNA
NM_022098	Homo sapiens hypothetical protein LOC63929 (LOC63929), mRNA
NM_022081	Homo sapiens hypothetical protein bK1048E9.5 (BK1048E9.5), mRNA
NM_021081	Homo sapiens growth hormone releasing hormone (GHRH), mRNA
NM_022168	Homo sapiens melanoma differentiation associated protein-5 (MDA5), mRNA
NM_022165	Homo sapiens Lin-7b protein (LIN-7B), mRNA
NM_022161	Homo sapiens livin inhibitor-of-apotosis (LIVIN), mRNA
NM_022159	Homo sapiens ETL protein (ETL), mRNA
NM_022156	Homo sapiens PP3111 protein (PP3111), mRNA
NM_022151	Homo sapiens MAP-1 protein (MAP-1), mRNA
NM_022150	Homo sapiens RFamide-related peptide precursor (RFRP), mRNA
NM_022149	Homo sapiens MAGEF1 protein (MAGEF1), mRNA
NM_022144	Homo sapiens myodulin protein (LOC64102), mRNA
NM_022141	Homo sapiens gamma-parvin (PARVG), mRNA
NM_022134	Homo sapiens glycoprotein beta-Gal 3'-sulfotransferase (GP3ST), mRNA
NM_022131	Homo sapiens calsyntenin-2 (CS2), mRNA
NM_022129	Homo sapiens MAWD binding protein (MAWBP), mRNA
NM_022123	Homo sapiens basic-helix-loop-helix-PAS protein (NPAS3), mRNA
NM_022121	Homo sapiens p53-induced protein PIGPC1 (PIGPC1), mRNA
NM_022120	Homo sapiens hypothetical protein FKSG25 (FLJ00030), mRNA
NM_022114	Homo sapiens PR domain containing 16 (PRDM16), mRNA
NM_022112	Homo sapiens p53-regulated apoptosis-inducing protein 1 (P53AIP1), mRNA
NM_022111	Homo sapiens homolog of Xenopus Claspin (CLASPIN), mRNA
NM_022101	Homo sapiens hypothetical protein FLJ22965 (FLJ22965), mRNA
NM_022087	Homo sapiens hypothetical protein FLJ21634 (FLJ21634), mRNA
NM_022083	Homo sapiens niban protein (NIBAN), mRNA
NM_022078	Homo sapiens hypothetical protein FLJ12455 (FLJ12455), mRNA
NM_022076	Homo sapiens hypothetical protein IMAGE 109914 (LOC63904), mRNA
NM_022072	Homo sapiens hypothetical protein FLJ22609 (FLJ22609), mRNA
NM_022067	Homo sapiens hypothetical protein FLJ12707 (FLJ12707), mRNA
NM_022049	Homo sapiens G-protein coupled receptor 88 (GPR88), mRNA
NM_022044	Homo sapiens stromal cell-derived factor 2-like 1 (SDF2L1), mRNA
NM_022042	Homo sapiens solute carrier family 26 (sulfate transporter), member 1
-	(SLC26A1), mRNA
NM 022039	Homo sapiens split hand/foot malformation (ectrodactyly) type 3 (SHFM3),
-	mRNA
,	
NM_021173	Homo sapiens polymerase (DNA-directed), delta 4 (POLD4), mRNA

NM_000023	Homo sapiens sarcoglycan, alpha (50kD dystrophin-associated glycoprotein) (SGCA), mRNA
NM_005099	Homo sapiens a disintegrin-like and metalloprotease (reprolysin type) with
NR 016500	thrombospondin type 1 motif, 4 (ADAMTS4), mRNA
NM_016590	Homo sapiens prostate androgen-regulated transcript 1 (PART1), mRNA
NM_014223	Homo sapiens nuclear transcription factor Y, gamma (NFYC), mRNA
NM_006166	Homo sapiens nuclear transcription factor Y, beta (NFYB), mRNA
NM_002268	Homo sapiens karyopherin alpha 4 (importin alpha 3) (KPNA4), mRNA
NM_005229	Homo sapiens ELK1, member of ETS oncogene family (ELK1), mRNA
NM_021796 NM_015596	Homo sapiens placenta-specific 1 (PLAC1), mRNA  Homo sapiens kallikrein 13 (KLK13), mRNA
	Homo sapiens olfactory receptor, family 1, subfamily E, member 1 (OR1E1),
NM_003553	mRNA
NM_021926	Homo sapiens aristaless-like homeobox 4 (ALX4), mRNA
NM_021957	Homo sapiens glycogen synthase 2 (liver) (GYS2), mRNA
NM_020980	Homo sapiens aquaporin 9 (AQP9), mRNA
NM_001614	Homo sapiens actin, gamma 1 (ACTG1), mRNA
NM_018690	Homo sapiens apolipoprotein B48 receptor (APOB48R), mRNA
NM_005230	Homo sapiens ELK3, ETS-domain protein (SRF accessory protein 2) (ELK3), mRNA
NM_003816	Homo sapiens a disintegrin and metalloproteinase domain 9 (meltrin gamma) (ADAM9), mRNA
NM 000847	Homo sapiens glutathione S-transferase A3 (GSTA3), mRNA
NM_021814	Homo sapiens homolog of yeast long chain polyunsaturated fatty acid elongation enzyme 2 (HELO1), mRNA
NM 021628	Homo sapiens arachidonate lipoxygenase 3 (ALOXE3), mRNA
NM 012419	Homo sapiens regulator of G-protein signalling 17 (RGS17), mRNA
NM_014685	Homo sapiens homocysteine-inducible, endoplasmic reticulum stress-inducible, ubiquitin-like domain member 1 (HERPUD1), mRNA
NM 005705	Homo sapiens pan-hematopoietic expression (PHEMX), mRNA
NM 004906	Homo sapiens Wilms' tumour 1-associating protein (KIAA0105), mRNA
NM_003101	Homo sapiens sterol O-acyltransferase (acyl-Coenzyme A cholesterol acyltransferase) 1 (SOAT1), mRNA
NM 021965	Homo sapiens phosphoglucomutase 5 (PGM5), mRNA
NM_003555	Homo sapiens olfactory receptor, family 1, subfamily G, member 1 (OR1G1),
NM_003552	mRNA Homo sapiens olfactory receptor, family 1, subfamily D, member 4 (OR1D4),
NIM 001245	mRNA Homo sapiens diacylglycerol kinase, alpha (80kD) (DGKA), mRNA
NM_001345	<u> </u>
NM 021620 NM 020999	Homo sapiens PR domain containing 13 (PRDM13), mRNA Homo sapiens neurogenin 3 (NEUROG3), mRNA
NM 020227	Homo sapiens PR domain containing 9 (PRDM9), mRNA
NM 020226	Homo sapiens PR domain containing 9 (PRDM9), mRNA  Homo sapiens PR domain containing 8 (PRDM8), mRNA
NM 020229	Homo sapiens PR domain containing 3 (PRDM1), mRNA  Homo sapiens PR domain containing 11 (PRDM11), mRNA
NM 020228	Homo sapiens PR domain containing 11 (PRDM11), mRNA  Homo sapiens PR domain containing 10 (PRDM10), mRNA
NM 016412	Homo sapiens insulin-like growth factor 2, antisense (IGF2AS), mRNA
NM 006161	Homo sapiens neurogenin 1 (NEUROG1), mRNA
NM 005734	Homo sapiens homeodomain-interacting protein kinase 3 (HIPK3), mRNA
NM 001818	Homo sapiens aldo-keto reductase family 1, member C4 (chlordecone reductase;
14141_001010	3-alpha hydroxysteroid dehydrogenase, type I; dihydrodiol dehydrogenase 4) (AKR1C4), mRNA
NM 004363	Homo sapiens carcinoembryonic antigen-related cell adhesion molecule 5

	(CEACAM5), mRNA
NM 002841	Homo sapiens protein tyrosine phosphatase, receptor type, G (PTPRG), mRNA
NM 002716	Homo sapiens protein phosphatase 2 (formerly 2A), regulatory subunit A (PR
NWI_002716	65), beta isoform (PPP2R1B), mRNA
NM 001785	Homo sapiens cytidine deaminase (CDA), mRNA
NM 003554	Homo sapiens olfactory receptor, family 1, subfamily E, member 2 (OR1E2),
14141_005554	mRNA
NM 021961	Homo sapiens TEA domain family member 1 (SV40 transcriptional enhancer
	factor) (TEAD1), mRNA
NM 002847	Homo sapiens protein tyrosine phosphatase, receptor type, N polypeptide 2
	(PTPRN2), mRNA
NM_002778	Homo sapiens prosaposin (variant Gaucher disease and variant metachromatic
	leukodystrophy) (PSAP), mRNA
NM_000934	Homo sapiens serine (or cysteine) proteinase inhibitor, clade F (alpha-2
_	antiplasmin, pigment epithelium derived factor), member 2 (SERPINF2), mRNA
NM_000932	Homo sapiens phospholipase C, beta 3 (phosphatidylinositol-specific) (PLCB3),
	mRNA
NM_000709	Homo sapiens branched chain keto acid dehydrogenase E1, alpha polypeptide
	(maple syrup urine disease) (BCKDHA), mRNA
NM_001666	Homo sapiens Rho GTPase activating protein 4 (ARHGAP4), mRNA
NM_021815	Homo sapiens solute carrier family 5 (choline transporter), member 7 (SLC5A7),
	mRNA 10 (4 PG10) PNA
NM_014885	Homo sapiens anaphase-promoting complex 10 (APC10), mRNA
NM_021948	Homo sapiens chondroitin sulfate proteoglycan BEHAB/brevican (BCAN),
	mRNA
NM_021946	Homo sapiens hypothetical protein FLJ11362 (FLJ11362), mRNA
NM_021942	Homo sapiens hypothetical protein FLJ12716 (FLJ12716), mRNA
NM_021940	Homo sapiens hypothetical protein FLJ13159 (FLJ13159), mRNA Homo sapiens Fanconi anemia, complementation group E (FANCE), mRNA
NM 021922	Homo sapiens polymeric immunoglobulin receptor (PIGR), mRNA
NM_002644	Homo sapiens myosin, heavy polypeptide 3, skeletal muscle, embryonic
NM_002470	(MYH3), mRNA
NM 001700	Homo sapiens azurocidin 1 (cationic antimicrobial protein 37) (AZU1), mRNA
NM 003949	Homo sapiens huntingtin-associated protein 1 (neuroan 1) (HAP1), mRNA
NM 021021	Homo sapiens syntrophin, beta 1 (dystrophin-associated protein A1, 59kD, basic
14141_021021	component 1) (SNTB1), mRNA
NM 018953	Homo sapiens homeo box C5 (HOXC5), mRNA
NM 012120	Homo sapiens CD2-associated protein (CD2AP), mRNA
NM 007121	Homo sapiens nuclear receptor subfamily 1, group H, member 2 (NR1H2),
	mRNA
NM 006753	Homo sapiens surfeit 6 (SURF6), mRNA
NM 006200	Homo sapiens proprotein convertase subtilisin/kexin type 5 (PCSK5), mRNA
NM 006426	Homo sapiens dihydropyrimidinase-like 4 (DPYSL4), mRNA
NM_005670	Homo sapiens epilepsy, progressive myoclonus type 2, Lafora disease (laforin)
	(EPM2A), mRNA
NM_006877	Homo sapiens guanosine monophosphate reductase (GMPR), mRNA
NM_004619	Homo sapiens TNF receptor-associated factor 5 (TRAF5), mRNA
NM_002627	Homo sapiens phosphofructokinase, platelet (PFKP), mRNA
NM_002433	Homo sapiens myelin oligodendrocyte glycoprotein (MOG), mRNA
NM_002207	Homo sapiens integin, alpha 9 (ITGA9), mRNA
NM_002113	Homo sapiens H factor (complement)-like 1 (HFL1), mRNA
NM_002074	Homo sapiens guanine nucleotide binding protein (G protein), beta polypeptide 1

	(GNB1), mRNA
NM 003733	Homo sapiens 2'-5'oligoadenylate synthetase-like (OASL), mRNA
NM 002551	Homo sapiens olfactory receptor, family 3, subfamily A, member 2 (OR3A2),
NMI_002551	mRNA
NM_002389	Homo sapiens membrane cofactor protein (CD46, trophoblast-lymphocyte cross-
	reactive antigen) (MCP), mRNA
NM_000870	Homo sapiens 5-hydroxytryptamine (serotonin) receptor 4 (HTR4), mRNA
NM_000613	Homo sapiens hemopexin (HPX), mRNA
NM_000377	Homo sapiens Wiskott-Aldrich syndrome (eczema-thrombocytopenia) (WAS), mRNA
NM_006981	Homo sapiens nuclear receptor subfamily 4, group A, member 3 (NR4A3), mRNA
NM 000368	Homo sapiens TSC1 gene (hamartin) (TSC1), mRNA
NM_017416	Homo sapiens interleukin 1 receptor accessory protein-like 2 (IL1RAPL2), mRNA
NM 003286	Homo sapiens topoisomerase (DNA) I (TOP1), mRNA
NM 001068	Homo sapiens topoisomerase (DNA) II beta (180kD) (TOP2B), mRNA
NM 020470	Homo sapiens putative transmembrane protein; homolog of yeast Golgi
	membrane protein Yiflp (Yiplp-interacting factor) (54TM), mRNA
NM_006562	Homo sapiens transcription factor similar to D. melanogaster homeodomain protein lady bird late (LBX1), mRNA
NM 017545	Homo sapiens hydroxyacid oxidase (glycolate oxidase) 1 (HAO1), mRNA
NM 002925	Homo sapiens regulator of G-protein signalling 10 (RGS10), mRNA
NM 012263	Homo sapiens tubulin tyrosine ligase-like 1 (TTLL1), mRNA
NM 001212	Homo sapiens complement component 1, q subcomponent binding protein
11112_001_11	(C1QBP), nuclear gene encoding mitochondrial protein, mRNA
NM_000491	Homo sapiens complement component 1, q subcomponent, beta polypeptide (C1QB), mRNA
NM_004720	Homo sapiens endothelial differentiation, lysophosphatidic acid G-protein- coupled receptor, 4 (EDG4), mRNA
NM_006217	Homo sapiens serine (or cysteine) proteinase inhibitor, clade I (neuroserpin), member 2 (SERPINI2), mRNA
NM 018723	Homo sapiens ataxin 2-binding protein 1 (A2BP1), mRNA
NM 004543	Homo sapiens nebulin (NEB), mRNA
NM 016151	Homo sapiens prostate derived STE20-like kinase PSK (PSK), mRNA
NM 016528	Homo sapiens hydroxyacid oxidase 3 (medium-chain) (HAO3), mRNA
NM_000185	Homo sapiens serine (or cysteine) proteinase inhibitor, clade D (heparin cofactor), member 1 (SERPIND1), mRNA
NM 005410	Homo sapiens selenoprotein P, plasma, 1 (SEPP1), mRNA
NM_005226	Homo sapiens endothelial differentiation, sphingolipid G-protein-coupled receptor, 3 (EDG3), mRNA
NM 005172	Homo sapiens atonal homolog 1 (Drosophila) (ATOH1), mRNA
NM 005109	Homo sapiens oxidative-stress responsive 1 (OSR1), mRNA
NM 001498	Homo sapiens glutamate-cysteine ligase, catalytic subunit (GCLC), mRNA
NM 003922	Homo sapiens heet (homologous to the E6-AP (UBE3A) carboxyl terminus)
1444_005722	domain and RCC1 (CHC1)-like domain (RLD) 1 (HERC1), mRNA
NM 002061	Homo sapiens glutamate-cysteine ligase, modifier subunit (GCLM), mRNA
NM 001088	Homo sapiens arylalkylamine N-acetyltransferase (AANAT), mRNA
NM_021828	Homo sapiens heparanase-like protein (HPA2), mRNA
NM 021826	Homo sapiens hypothetical protein FLJ13149 (FLJ13149), mRNA
NM 021823	Homo sapiens hypothetical protein MDS018 (MDS018), mRNA
NM 021820	Homo sapiens MDS024 protein (MDS024), mRNA
1111 021020	1 Home suprems removed protein (removed), meeta

	TO CT (TID CT) DATA
NM_021819	Homo sapiens ERGL protein (ERGL), mRNA
NM_021818	Homo sapiens WW Domain-Containing Gene (WW45), mRNA
NM_021812	Homo sapiens blepharophimosis, epicanthus inversus and ptosis, candidate 1
	(BPESC1), mRNA
NM_021809	Homo sapiens TGF(beta)-induced transcription factor 2 (TGIF2), mRNA
NM_021805	Homo sapiens single Ig IL-1R-related molecule (SIGIRR), mRNA
NM_021803	Homo sapiens interleukin 21 (IL21), mRNA
NM_021798	Homo sapiens interleukin 21 receptor (IL21R), mRNA
NM_020982	Homo sapiens claudin 9 (CLDN9), mRNA
NM 006657	Homo sapiens formiminotransferase cyclodeaminase (FTCD), mRNA
NM 021784	Homo sapiens hepatocyte nuclear factor 3, beta (HNF3B), mRNA
NM 014375	Homo sapiens fetuin B (FETUB), mRNA
NM 021032	Homo sapiens fibroblast growth factor 12 (FGF12), mRNA
NM 019595	Homo sapiens intersectin 2 (ITSN2), mRNA
NM 018991	Homo sapiens DKFZp434A0131 protein (DKFZP434A0131), mRNA
NM 014574	Homo sapiens nuclear autoantigen (GS2NA), mRNA
NM 021002	Homo sapiens interferon, alpha 6 (IFNA6), mRNA
NM 001676	Homo sapiens ATPase, H+/K+ transporting, nongastric, alpha polypeptide
_	(ATP12A), mRNA
NM_019886	Homo sapiens carbohydrate (N-acetylglucosamine 6-O) sulfotransferase 7
	(CHST7), mRNA
NM 017581	Homo sapiens cholinergic receptor, nicotinic, alpha polypeptide 9 (CHRNA9),
	mRNA
NM 001695	Homo sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump)
	42kD (ATP6C), mRNA
NM 006303	Homo sapiens JTV1 gene (JTV1), mRNA
NM 014413	Homo sapiens heme-regulated initiation factor 2-alpha kinase (HRI), mRNA
NM 012149	Homo sapiens double homeobox, 5 (DUX5), mRNA
NM_012146	Homo sapiens double homeobox, 1 (DUX1), mRNA
NM 021733	Homo sapiens testis-specific kinase substrate (TSKS), mRNA
NM 004339	Homo sapiens pituitary tumor-transforming 1 interacting protein (PTTG1IP),
_	mRNA
NM 004219	Homo sapiens pituitary tumor-transforming 1 (PTTG1), mRNA
NM 003860	Homo sapiens Breakpoint cluster region protein, uterine leiomyoma, 1; barrier to
_	autointegration factor (BCRP1), mRNA
NM 007281	Homo sapiens scrapie responsive protein 1 (SCRG1), mRNA
NM 006618	Homo sapiens putative DNA/chromatin binding motif (PLU-1), mRNA
NM 005797	Homo sapiens epithelial V-like antigen 1 (EVA1), mRNA
NM 005508	Homo sapiens chemokine (C-C motif) receptor 4 (CCR4), mRNA
NM 005283	Homo sapiens chemokine (C motif) XC receptor 1 (CCXCR1), mRNA
NM 002547	Homo sapiens oligophrenin 1 (OPHN1), mRNA
NM 020056	Homo sapiens major histocompatibility complex, class II, DQ alpha 2 (HLA-
	DQA2), mRNA
NM_001085	Homo sapiens serine (or cysteine) proteinase inhibitor, clade A (alpha-1
_	antiproteinase, antitrypsin), member 3 (SERPINA3), mRNA
NM 013974	Homo sapiens dimethylarginine dimethylaminohydrolase 2 (DDAH2), mRNA
NM 001756	Homo sapiens serine (or cysteine) proteinase inhibitor, clade A (alpha-1
	antiproteinase, antitrypsin), member 6 (SERPINA6), mRNA
NM_000450	Homo sapiens selectin E (endothelial adhesion molecule 1) (SELE), mRNA
NM 006228	Homo sapiens prepronociceptin (PNOC), mRNA
NM 001319	Homo sapiens casein kinase 1, gamma 2 (CSNK1G2), mRNA
NM 000444	Homo sapiens phosphate regulating gene with homologies to endopeptidases on

the X chromosome (hypophosphatemia, vitamin D resistant rickets) (PHEX),
mRNA
Homo sapiens hypothetical protein similar to small G proteins, especially RAP-
2A (LOC57826), mRNA
Homo sapiens hypothetical protein LOC57821 (LOC57821), mRNA
Homo sapiens protein kinase C, zeta (PRKCZ), mRNA
Homo sapiens serine (or cysteine) proteinase inhibitor, clade A (alpha-1
antiproteinase, antitrypsin), member 5 (SERPINA5), mRNA
Homo sapiens serine (or cysteine) proteinase inhibitor, clade E (nexin,
plasminogen activator inhibitor type 1), member 1 (SERPINE1), mRNA
Homo sapiens hypothetical protein from clone 24796 (LOC57146), mRNA
Homo sapiens transcription factor BMAL2 (LOC56938), mRNA
Homo sapiens kallikrein 12 (KLK12), mRNA
Homo sapiens hypothetical protein (LOC55954), mRNA
Homo sapiens serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 13 (SERPINB13), mRNA
Homo sapiens low density lipoprotein receptor (familial hypercholesterolemia)
(LDLR), mRNA
Homo sapiens U6 snRNA-associated Sm-like protein LSm8 (LOC51691), mRNA
Homo sapiens KIAA0193 gene product (KIAA0193), mRNA
Homo sapiens RNA binding motif protein 9 (RBM9), mRNA
Homo sapiens dual oxidase-like domains 2 (DUOX2), mRNA
Homo sapiens CCR4-NOT transcription complex, subunit 3 (CNOT3), mRNA
Homo sapiens KIAA0979 protein (KIAA0979), mRNA
Homo sapiens KIAA0040 gene product (KIAA0040), mRNA
Homo sapiens hypothetical protein (DJ328E19.C1.1), mRNA
Homo sapiens protein kinase C, alpha binding protein (PRKCABP), mRNA
Homo sapiens integrin, alpha E (antigen CD103, human mucosal lymphocyte antigen 1; alpha polypeptide) (ITGAE), mRNA
Homo sapiens leukemia inhibitory factor (cholinergic differentiation factor) (LIF), mRNA
Homo sapiens serine (or cysteine) proteinase inhibitor, clade B (ovalbumin),
member 3 (SERPINB3), mRNA
Homo sapiens serine (or cysteine) proteinase inhibitor, clade A (alpha-1
antiproteinase, antitrypsin), member 2 (SERPINA2), mRNA
Homo sapiens serine (or cysteine) proteinase inhibitor, clade A (alpha-1
antiproteinase, antitrypsin), member 4 (SERPINA4), mRNA
Homo sapiens deleted in lymphocytic leukemia, 2 (DLEU2), mRNA
Homo sapiens deleted in lymphocytic leukemia, 1 (DLEU1), mRNA
Homo sapiens ATPase, Class I, type 8B, member 1 (ATP8B1), mRNA
Homo sapiens EphA1 (EPHA1), mRNA
Homo sapiens serine (or cysteine) proteinase inhibitor, clade B (ovalbumin),
member 10 (SERPINB10), mRNA
Homo sapiens CCR4-NOT transcription complex, subunit 8 (CNOT8), mRNA
Homo sapiens serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 9 (SERPINB9), mRNA
Homo sapiens serine (or cysteine) proteinase inhibitor, clade B (ovalbumin),
Homo sapiens serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 6 (SERPINB6), mRNA
Homo sapiens serine (or cysteine) proteinase inhibitor, clade B (ovalbumin),

NM_000214	Homo sapiens jagged 1 (Alagille syndrome) (JAG1), mRNA
NM 001347	Homo sapiens diacylglycerol kinase, theta (110kD) (DGKQ), mRNA
NM 003454	Homo sapiens zinc finger protein 200 (ZNF200), mRNA
NM 003334	Homo sapiens ubiquitin-activating enzyme E1 (A1S9T and BN75 temperature
·	sensitivity complementing) (UBE1), mRNA
NM_000354	Homo sapiens serine (or cysteine) proteinase inhibitor, clade A (alpha-1
	antiproteinase, antitrypsin), member 7 (SERPINA7), mRNA
NM_000945	Homo sapiens protein phosphatase 3 (formerly 2B), regulatory subunit B (19kD),
11111_0005.5	alpha isoform (calcineurin B, type I) (PPP3R1), mRNA
NM 000305	Homo sapiens paraoxonase 2 (PON2), mRNA
NM 000928	Homo sapiens phospholipase A2, group IB (pancreas) (PLA2G1B), nuclear gene
14141_000520	encoding mitochondrial protein, mRNA
NM_000295	Homo sapiens serine (or cysteine) proteinase inhibitor, clade A (alpha-1
14141_000293	antiproteinase, antitrypsin), member 1 (SERPINA1), mRNA
NM 002640	Homo sapiens serine (or cysteine) proteinase inhibitor, clade B (ovalbumin),
NWI_002040	member 8 (SERPINB8), mRNA
NTM 002620	Homo sapiens serine (or cysteine) proteinase inhibitor, clade B (ovalbumin),
NM_002639	member 5 (SERPINB5), mRNA
2D 6 000615	Homo sapiens serine (or cysteine) proteinase inhibitor, clade F (alpha-2
NM_002615	antiplasmin, pigment epithelium derived factor), member 1 (SERPINF1), mRNA
ND 6 000575	Homo sapiens serine (or cysteine) proteinase inhibitor, clade B (ovalbumin),
NM_002575	Homo sapiens serine (or cysteine) proteinase infinition, clade is (ovaloumin),
37.5.000000	member 2 (SERPINB2), mRNA  Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 1
NM_000220	
	(KCNJ1), mRNA
NM_000191	Homo sapiens 3-hydroxymethyl-3-methylglutaryl-Coenzyme A lyase
	(hydroxymethylglutaricaciduria) (HMGCL), mRNA
NM_001978	Homo sapiens erythrocyte membrane protein band 4.9 (dematin) (EPB49),
	mRNA
NM_003646	Homo sapiens diacylglycerol kinase, zeta (104kD) (DGKZ), mRNA
NM_001346	Homo sapiens diacylglycerol kinase, gamma (90kD) (DGKG), mRNA
NM_003647	Homo sapiens diacylglycerol kinase, epsilon (64kD) (DGKE), mRNA
NM_001235	Homo sapiens serine (or cysteine) proteinase inhibitor, clade H (heat shock
	protein 47), member 2 (SERPINH2), mRNA
NM_001694	Homo sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump)
	16kD (ATP6L), mRNA
NM_000488	Homo sapiens serine (or cysteine) proteinase inhibitor, clade C (antithrombin),
	member 1 (SERPINC1), mRNA
NM_021156	Homo sapiens hypothetical protein (DJ971N18.2), mRNA
NM_000875	Homo sapiens insulin-like growth factor 1 receptor (IGF1R), mRNA
NM_000605	Homo sapiens interferon, alpha 2 (IFNA2), mRNA
NM_021647	Homo sapiens KIAA0626 gene product (KIAA0626), mRNA
NM_021645	Homo sapiens KIAA0266 gene product (KIAA0266), mRNA
NM_021109	Homo sapiens thymosin, beta 4, X chromosome (TMSB4X), mRNA
NM_021642	Homo sapiens Fc fragment of IgG, low affinity IIa, receptor for (CD32)
_	(FCGR2A), mRNA
NM 021240	Homo sapiens testis-specific protein (LOC58524), mRNA
NM 021189	Homo sapiens hypothetical protein FLJ10698 (LOC57863), mRNA
NM 021129	Homo sapiens pyrophosphatase (inorganic) (PP), nuclear gene encoding
	mitochondrial protein, mRNA
NM 015140	Homo sapiens KIAA0153 protein (KIAA0153), mRNA
NM 021635	Homo sapiens UC28 protein (UC28), mRNA
NM 021631	Homo sapiens apoptosis inhibitor (FKSG2), mRNA
14141 07 102 1	Tromo suprema apopusta intitotion (***1502); ****

NM_021615	
	Homo sapiens carbohydrate (N-acetylglucosamine 6-O) sulfotransferase 6 (CHST6), mRNA
NM 012334	Homo sapiens myosin X (MYO10), mRNA
NM 020363	Homo sapiens deleted in azoospermia 2 (DAZ2), mRNA
NM 020364	Homo sapiens deleted in azoospermia 3 (DAZ3), mRNA
NM 017445	Homo sapiens H2B histone family, member S (H2BFS), mRNA
NM 021132	Homo sapiens protein phosphatase 3 (formerly 2B), catalytic subunit, beta
_	isoform (calcineurin A beta) (PPP3CB), mRNA
NM 021016	Homo sapiens pregnancy specific beta-1-glycoprotein 3 (PSG3), mRNA
NM 015705	Homo sapiens hypothetical protein (DJ1042K10.2), mRNA
NM_021572	Homo sapiens ectonucleotide pyrophosphatase/phosphodiesterase 5 (putative
_	function) (ENPP5), mRNA
NM_021216	Homo sapiens endothelial zinc finger protein induced by tumor necrosis factor
-	alpha (EZFIT), mRNA
NM_001332	Homo sapiens catenin (cadherin-associated protein), delta 2 (neural plakophilin-
_	related arm-repeat protein) (CTNND2), mRNA
NM_021185	Homo sapiens hypothetical protein DKFZp434A1022 (DKFZP434A1022),
_	mRNA
NM 018955	Homo sapiens ubiquitin B (UBB), mRNA
NM 017533	Homo sapiens myosin, heavy polypeptide 4, skeletal muscle (MYH4), mRNA
NM 014621	Homo sapiens homeo box D4 (HOXD4), mRNA
NM 000618	Homo sapiens insulin-like growth factor 1 (somatomedia C) (IGF1), mRNA
NM 021571	Homo sapiens ICEBERG caspase-1 inhibitor (ICEBERG), mRNA
NM 000045	Homo sapiens arginase, liver (ARG1), mRNA
NM 005692	Homo sapiens ATP-binding cassette, sub-family F (GCN20), member 2
	(ABCF2), mRNA
NM_001090	Homo sapiens ATP-binding cassette, sub-family F (GCN20), member 1 (ABCF1), mRNA
NM_002858	Homo sapiens ATP-binding cassette, sub-family D (ALD), member 3 (ABCD3), mRNA
NM_001172	Homo sapiens arginase, type II (ARG2), nuclear gene encoding mitochondrial protein, mRNA
NM_001117	Homo sapiens adenylate cyclase activating polypeptide 1 (pituitary) (ADCYAP1), mRNA
NM 004036	Homo sapiens adenylate cyclase 3 (ADCY3), mRNA
NM 019843	Homo sapiens eIF4E-transporter (4E-T), mRNA
NM 006454	Homo sapiens Mad4 homolog (MAD4), mRNA
NM_002355	Homo sapiens mannose-6-phosphate receptor (cation dependent) (M6PR), mRNA
NM_014287	Homo sapiens pM5 protein (PM5), mRNA
NM_004102	Homo sapiens fatty acid binding protein 3, muscle and heart (mammary-derived growth inhibitor) (FABP3), mRNA
NM 000134	Homo sapiens fatty acid binding protein 2, intestinal (FABP2), mRNA
NM 005354	Homo sapiens jun D proto-oncogene (JUND), mRNA
NM_005159	Homo sapiens actin, alpha, cardiac muscle (ACTC), mRNA
NM 019848	Homo sapiens Protein P3 (P3), mRNA
NM_003948	Homo sapiens cyclin-dependent kinase-like 2 (CDC2-related kinase) (CDKL2), mRNA
NM_021131	Homo sapiens protein phosphatase 2A, regulatory subunit B' (PR 53) (PPP2R4), mRNA
1	<del></del>
NM 021268	Homo sapiens interferon, alpha 17 (IFNA17), mRNA

NM 001166 Homo sapiens X-prolyl aminopeptidase (aminopeptidase) Homo sapiens X-prolyl aminopeptidase (aminopeptidase) P. 2, membrane-bound (XPNPEP2), mRNA  NM 005319 Homo sapiens smosin regulatory light chain interacting protein (MIR), mRNA  NM 005393 Homo sapiens plexin B3 (PLXNB3), mRNA  NM 005393 Homo sapiens plexim B3 (PLXNB3), mRNA  NM 021287 Homo sapiens neuroglobin (NGB), mRNA  NM 021257 Homo sapiens ring finger protein 23 (RNF23), mRNA  NM 021247 Homo sapiens protamine 3 (PRM3), mRNA  NM 021247 Homo sapiens protamine 3 (PRM3), mRNA  NM 021242 Homo sapiens ring finger protein 3 (RNF23), mRNA  NM 021242 Homo sapiens Protein (TERA), mRNA  NM 021223 Homo sapiens TERA protein (TERA), mRNA  NM 021221 Homo sapiens Stroptotin (GSB), mRNA  NM 021221 Homo sapiens Stroptotin (GSB), mRNA  NM 021220 Homo sapiens EST-YD1 protein (MUM2), mRNA  NM 021200 Homo sapiens EST-YD1 protein (MUM2), mRNA  NM 021200 Homo sapiens EST-YD1 protein (EST-YD1), mRNA  NM 021100 Homo sapiens EGT-44 protein; sulfide dehydrogenase like (yeast) (CGI-44), mRNA  NM 021198 Homo sapiens muclear LIM interactor-interacting factor (NLI-IF), mRNA  NM 021198 Homo sapiens bomeo box D12 (HOXD12), mRNA  NM 021194 Homo sapiens bomeo box D11 (HOXD11), mRNA  NM 021195 Homo sapiens D05 (POXD12), mRNA  NM 021184 Homo sapiens Potentin (GNXD11), mRNA  NM 021185 Homo sapiens bomeo box D12 (HOXD12), mRNA  NM 021186 Homo sapiens Potentin (GNXD11), mRNA  NM 021187 Homo sapiens Potentin (GNXD11), mRNA  NM 021187 Homo sapiens Surve to sapiens Surv		The secondary of the se
(XPNPEP2), mRNA  NM 00393 Homo sapiens S-antigen; retina and pineal gland (arrestin) (SAG), mRNA NM 013262 Homo sapiens myosin regulatory light chain interacting protein (MIR), mRNA NM 00393 Homo sapiens plexin B3 (PLXNB3), mRNA NM 021257 Homo sapiens calcium channel, voltage-dependent, alpha 1H subunit (CACNA1H), mRNA NM 021257 Homo sapiens rium finger protein 23 (RNF23), mRNA NM 021247 Homo sapiens protamine 3 (PRM3), mRNA NM 021242 Homo sapiens protamine 3 (PRM3), mRNA NM 021242 Homo sapiens brothetical protein STRAIT 11499 (STRAIT11499), mRNA NM 021243 Homo sapiens TERA protein (TERA), mRNA NM 021221 Homo sapiens Stpotnetical protein STRAIT 11499 (STRAIT11499), mRNA NM 021221 Homo sapiens Stpotnetin (GSB), mRNA NM 021221 Homo sapiens Stpotnetin (GSB), mRNA NM 021201 Homo sapiens BUM2 protein (MUM2), mRNA NM 021201 Homo sapiens BUM2 protein (MUM2), mRNA NM 021200 Homo sapiens EST-YD1 protein (EST-YD1), mRNA NM 021109 Homo sapiens CGI-44 protein; sulfide dehydrogenase like (yeast) (CGI-44), mRNA NM 021199 Homo sapiens bome box D11 (HOXD11), mRNA NM 021191 Homo sapiens Sen borneo box D11 (HOXD11), mRNA NM 021192 Homo sapiens shorneo box D11 (HOXD11), mRNA NM 021193 Homo sapiens shorneo box D11 (HOXD11), mRNA NM 021194 Homo sapiens Sen borneo box D11 (HOXD11), mRNA NM 021174 Homo sapiens G4 protein (G4), mRNA NM 021175 Homo sapiens Sen spiens brothe obx D11 (MOXD11), mRNA NM 021174 Homo sapiens Sen protein (G4), mRNA NM 021175 Homo sapiens Sen protein (G4), mRNA NM 021176 Homo sapiens Sen protein (G4), mRNA NM 021177 Homo sapiens Sen protein (G4), mRNA NM 021177 Homo sapiens Sen protein (G4), mRNA NM 021177 Homo sapiens Sen protein (G4), mRNA NM 021174 Homo sapiens Sen protein (G4), mRNA NM 021175 Homo sapiens Sen protein (G4), mRNA NM 021176 Homo sapiens Sen protein (G4), mRNA NM 021177 Homo sapiens Sen protein (G4), mRNA NM 021178 Homo sapiens Sen protein (G4), mRNA NM 021179 Homo sapiens Sen protein (G4), mRNA NM 021130 Homo sapiens Sen protein (G4), mRNA NM 021131 Homo sapiens Sen protein (G4), mRNA NM 0211	NM_001166	Homo sapiens baculoviral IAP repeat-containing 2 (BIRC2), mRNA
NM 000541   Homo sapiens S-antigen; retina and pineal gland (arrestin) (SAG), mRNA   NM 01362   Homo sapiens myosin regulatory light chain interacting protein (MIR), mRNA   NM 001393   Homo sapiens plexin B3 (PLXNB3), mRNA   NM 021098   Homo sapiens calcium channel, voltage-dependent, alpha 1H subunit (CACNA1H), mRNA   Homo sapiens neuroglobin (NGB), mRNA   NM 021257   Homo sapiens protein 23 (RNF23), mRNA   NM 021247   Homo sapiens protein 23 (RNF23), mRNA   NM 021247   Homo sapiens protein me 3 (PRM3), mRNA   NM 021247   Homo sapiens hypothetical protein STRAIT11499 (STRAIT11499), mRNA   NM 021223   Homo sapiens SGb protein (GERA), mRNA   NM 021221   Homo sapiens Gb protein (GERA), mRNA   NM 021221   Homo sapiens Gb protein (GSB), mRNA   NM 021208   Homo sapiens SGb protein (GSB), mRNA   NM 02100   Homo sapiens ET-YD1 protein (BST-YD1), mRNA   NM 02109   Homo sapiens CGI-44 protein; sulfide dehydrogenase like (yeast) (CGI-44), mRNA   NM 021193   Homo sapiens bulear LIM interactor-interacting factor (NLI-IF), mRNA   NM 021192   Homo sapiens box D12 (HOXD12), mRNA   NM 021193   Homo sapiens box D0x D12 (HOXD12), mRNA   NM 021184   Homo sapiens box D0x D12 (HOXD11), mRNA   NM 021184   Homo sapiens box D0x D12 (HOXD11), mRNA   NM 021184   Homo sapiens box D0x D12 (HOXD11), mRNA   NM 021184   Homo sapiens box D0x D12 (HOXD11), mRNA   NM 021184   Homo sapiens box D0x D12 (HOXD11), mRNA   NM 021184   Homo sapiens D0x D12 (HOXD12), mRNA   NM 021184   Homo sapiens D0x D12 (HOXD12), mRNA   NM 021184   Homo sapiens D0x D12 (HOXD11), mRNA   NM 021185   Homo sapiens D0x D12 (HOXD12), mRNA   NM 021185   Homo sapiens D0x D12 (HOXD12), mRNA   NM 021167   Homo sapiens D0x D12 (HOXD12), mRNA   NM 021167   Homo sapiens D0x D12 (HOXD12), mRNA   NM 021137   Homo sapiens D0x D12 (HOXD12), mRNA   NM 021137   Homo sapiens D0x D12 (HOXD12), mRNA   NM 021137   Homo sapiens D0x D12 (HOXD12), mRNA   NM 021139   Homo sapiens D0x D12 (HOXD12), mRNA   NM 021130   Homo sapiens D0x D12 (HOXD12), mRNA   NM 021130   Homo sapiens D0x D12 (HOX	NM_003399	(YPNPFP2) mRNA
NM 013262   Homo sapiens myosin regulatory light chain interacting protein (MIR), mRNA	NM 000541	Homo sapiens S-antigen: retina and pineal gland (arrestin) (SAG), mRNA
NM 005393   Homo sapiens plexin B3 (PLXNB3), mRNA		Homo sapiens myosin regulatory light chain interacting protein (MIR), mRNA
NM 021257   Homo sapiens calcium channel, voltage-dependent, alpha 1H subunit (CACNA1H), mRNA		Homo saniens plevin B3 (PLXNB3), mRNA
(CACNAIH), mRNA  NM 021257 Homo sapiens neuroglobin (NGB), mRNA  NM 021247 Homo sapiens neuroglobin (NGB), mRNA  NM 021247 Homo sapiens protamine 3 (PRM3), mRNA  NM 021248 Homo sapiens protamine 3 (PRM3), mRNA  NM 021238 Homo sapiens TERA protein (TERA), mRNA  NM 021221 Homo sapiens myosin light chain 2a (LOC58498), mRNA  NM 021221 Homo sapiens MUM2 protein (MUM2), mRNA  NM 021210 Homo sapiens MUM2 protein (MUM2), mRNA  NM 021210 Homo sapiens EST-YD1 protein (EST-YD1), mRNA  NM 021200 Homo sapiens EST-YD1 protein (BST-YD1), mRNA  NM 021201 Homo sapiens EST-YD1 protein (BST-YD1), mRNA  NM 021209 Homo sapiens EGT-YD1 protein (BST-YD1), mRNA  NM 021199 Homo sapiens NUM2 protein; sulfide dehydrogenase like (yeast) (CGI-44), mRNA  NM 021198 Homo sapiens nuclear LIM interactor-interacting factor (NLI-IF), mRNA  NM 021193 Homo sapiens homeo box D12 (HOXD12), mRNA  NM 021184 Homo sapiens homeo box D11 (HOXD11), mRNA  NM 021185 Homo sapiens GA protein (G4), mRNA  NM 021174 Homo sapiens U6 snRNA-associated Sm-like protein (LOC57862), mRNA  NM 021174 Homo sapiens J00 DBC protein (UOC57805), mRNA  NM 021174 Homo sapiens Homeo sapiens Wigner (CD209), mRNA  NM 021175 Homo sapiens CD209 antigen (CD209), mRNA  NM 021167 Homo sapiens CD209 antigen (CD209), mRNA  NM 021179 Homo sapiens U6 snRNA-associated Sm-like protein (LSM2), mRNA  NM 021174 Homo sapiens U6 snRNA-associated Sm-like protein (LSM2), mRNA  NM 021174 Homo sapiens Wigner (CD209), mRNA  NM 021175 Homo sapiens tracii-DNA glycosylase 2 (UNG2), mRNA  NM 021187 Homo sapiens tracii-DNA glycosylase 2 (UNG2), mRNA  NM 021130 Homo sapiens unor necrosis factor, alpha-induced protein 1 (endothelial) (TNFAIP1), mRNA  NM 021131 Homo sapiens stumor necrosis factor, alpha-induced protein 1 (endothelial) (TNFAIP1), mRNA  NM 021130 Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA  NM 021131 Homo sapiens situmor necrosis factor, alpha-induced protein 1 (endothelial) (ULG3), mRNA  NM 021130 Homo sapiens from the protein s6 kinase, 90kD, polypeptide 2 (RP		Homo sapiens calcium channel, voltage-dependent, alpha 1H subunit
NM 021257   Homo sapiens neuroglobin (NGB), mRNA	14141_021000	(CACNA1H), mRNA
NM 021242 Homo sapiens ring finger protein 23 (RNF23), mRNA NM 021242 Homo sapiens protamine 3 (PRM3), mRNA NM 021238 Homo sapiens hypothetical protein STRATT11499 (STRAIT11499), mRNA NM 021231 Homo sapiens myosin light chain 2a (LOC58498), mRNA NM 021211 Homo sapiens MVM2 protein (G5B), mRNA NM 021210 Homo sapiens STPA protein (G5B), mRNA NM 021210 Homo sapiens EST-YD1 protein (BST-YD1), mRNA NM 021200 Homo sapiens EST-YD1 protein (BST-YD1), mRNA NM 021200 Homo sapiens EST-YD1 protein (BST-YD1), mRNA NM 021290 Homo sapiens PH domain containing protein in retina 1 (PHRET1), mRNA NM 021199 Homo sapiens nuclear LIM interactor-interacting factor (NLI-IF), mRNA NM 021193 Homo sapiens homeo box D12 (HOXD12), mRNA NM 021193 Homo sapiens homeo box D11 (HOXD11), mRNA NM 021184 Homo sapiens clones 23667 and 23775 zinc finger protein (LOC57862), mRNA NM 021184 Homo sapiens G4 protein (G4), mRNA NM 021184 Homo sapiens U5 snRNA-associated Sm-like protein (LSM2), mRNA NM 021177 Homo sapiens p30 DBC protein (LOC57805), mRNA NM 021167 Homo sapiens RAP1, GTP-GDP dissociation stimulator 1 (RAP1GDS1), mRNA NM 021155 Homo sapiens CD209 antigen (CD209), mRNA NM 021157 Homo sapiens uracil-DNA glycosylase 2 (UNG2), mRNA NM 021130 Homo sapiens uracil-DNA glycosylase 2 (UNG2), mRNA NM 021131 Homo sapiens rotein (DOC57805), mRNA NM 021131 Homo sapiens rotein (DOC57805), mRNA NM 021131 Homo sapiens uracil-DNA glycosylase 2 (UNG2), mRNA NM 021131 Homo sapiens rotein (DOC57805), mRNA NM 021131 Homo sapiens uracil-DNA glycosylase 2 (UNG2), mRNA NM 021131 Homo sapiens rotein (DOC57805), mRNA NM 021131 Homo sapiens rotein (DOC57805), mRNA NM 021131 Homo sapiens stumor necrosis factor, alpha-induced protein 1 (endothelial) (TNFAIP1), mRNA NM 021131 Homo sapiens rotein (DOC57805), mRNA NM 021133 Homo sapiens rotein (DOC57805), mRNA NM 021134 Homo sapiens rotein (DOC57805), mRNA NM 021135 Homo sapiens rotein (DOC57805), mRNA NM 021130 Homo sapiens rotein (DOC57805), mRNA NM 021131 Homo sapiens rotein (DOC57805), mRNA NM 021131 Homo sapiens rote	NM 021257	Homo sapiens neuroglobin (NGB), mRNA
NM 021247   Homo sapiens protamine 3 (PRM3), mRNA		Homo sapiens ring finger protein 23 (RNF23), mRNA
NM 021232 Homo sapiens hypothetical protein STRAIT11499 (STRAIT11499), mRNA NM 021223 Homo sapiens TERA protein (TERA), mRNA NM 021221 Homo sapiens myosin light chain 2a (LOC58498), mRNA NM 021201 Homo sapiens MUM2 protein (MUM2), mRNA NM 021202 Homo sapiens SET-YD1 protein (EST-YD1), mRNA NM 021200 Homo sapiens SET-YD1 protein (EST-YD1), mRNA NM 021200 Homo sapiens SET-YD1 protein (EST-YD1), mRNA NM 021200 Homo sapiens CGI-44 protein; sulfide dehydrogenase like (yeast) (CGI-44), mRNA NM 021199 Homo sapiens nuclear LIM interactor-interacting factor (NLI-IF), mRNA NM 021193 Homo sapiens homeo box D12 (HOXD12), mRNA NM 021193 Homo sapiens homeo box D11 (HOXD11), mRNA NM 021184 Homo sapiens homeo box D11 (HOXD11), mRNA NM 021188 Homo sapiens clones 23667 and 23775 zinc finger protein (LOC57862), mRNA NM 021184 Homo sapiens GP protein (G4), mRNA NM 021174 Homo sapiens GP protein (LOC57805), mRNA NM 021177 Homo sapiens B30 DBC protein (LOC57805), mRNA NM 021167 Homo sapiens RAP1, GTP-GDP dissociation stimulator 1 (RAP1GDS1), mRNA NM 021159 Homo sapiens RAP1, GTP-GDP dissociation stimulator 1 (RAP1GDS1), mRNA NM 021140 Homo sapiens uracil-DNA glycosylase 2 (UNG2), mRNA NM 021147 Homo sapiens uracil-DNA glycosylase 2 (UNG2), mRNA NM 021130 Homo sapiens uracil-DNA glycosylase 2 (UNG2), mRNA NM 021131 Homo sapiens suracil-DNA glycosylase 2 (UNG2), mRNA NM 021130 Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide B4 (UGT2B4), mRNA NM 021133 Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA NM 021130 Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA NM 021130 Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA NM 021130 Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA NM 021130 Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA NM 021130 Homo sapiens ribonuclease L (2',5'-oligoisoadenylate synthetase-dependent) (RNASEL), mRNA NM 02120 Homo sapiens thyroid hormone receptor inter		Homo saniens protamine 3 (PRM3), mRNA
NM 021221 Homo sapiens TERA protein (TERA), mRNA NM 021221 Homo sapiens myosin light chain 2a (LOC58498), mRNA NM 021221 Homo sapiens G5b protein (G5B), mRNA NM 021201 Homo sapiens MUM2 protein (MUM2), mRNA NM 021200 Homo sapiens EST-YD1 protein (EST-YD1), mRNA NM 021200 Homo sapiens PH domain containing protein in retina 1 (PHRET1), mRNA NM 021199 Homo sapiens nuclear LIM interactor-interacting factor (NLI-IF), mRNA NM 021193 Homo sapiens homeo box D12 (HOXD12), mRNA NM 021194 Homo sapiens homeo box D12 (HOXD12), mRNA NM 021195 Homo sapiens homeo box D11 (HOXD11), mRNA NM 021188 Homo sapiens clones 23667 and 23775 zinc finger protein (LOC57862), mRNA NM 021184 Homo sapiens G4 protein (G4), mRNA NM 021177 Homo sapiens U6 snRNA-associated Sm-like protein (LSM2), mRNA NM 021174 Homo sapiens pDBC protein (LOC57805), mRNA NM 021167 Homo sapiens Pyothetical protein WUGSC:H_RG083M05.2 (LOC57798), mRNA NM 021159 Homo sapiens RAP1, GTP-GDP dissociation stimulator 1 (RAP1GDS1), mRNA NM 021140 Homo sapiens BAP1, GTP-GDP dissociation stimulator 1 (RAP1GDS1), mRNA NM 021147 Homo sapiens usiquitously transcribed tetratricopeptide repeat gene, X chromosome (UTX), mRNA NM 021140 Homo sapiens usiquitously transcribed tetratricopeptide repeat gene, X chromosome (UTX), mRNA NM 021138 Homo sapiens TNF receptor-associated factor 2 (TRAF2), mRNA NM 021137 Homo sapiens reticulon 1 (RTN1), mRNA NM 021138 Homo sapiens reticulon 1 (RTN1), mRNA NM 021130 Homo sapiens reticulon 1 (RTN1), mRNA NM 021131 Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA NM 021131 Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA NM 021130 Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA NM 021131 Homo sapiens discs, large (Drosophila) homolog 3 (neuroendocrine-dlg) (DLG3), mRNA NM 004239 Homo sapiens thyroid hormone receptor interactor 12 (TRP12), mRNA NM 004239 Homo sapiens discs, large (Drosophila) homolog-associated protein 2		Homo sapiens hypothetical protein STRAIT11499 (STRAIT11499), mRNA
NM   021223   Homo sapiens myosin light chain 2a (LOC58498), mRNA   NM   021210   Homo sapiens G5b protein (G5B), mRNA   NM   021208   Homo sapiens MUM2 protein (MUM2), mRNA   NM   021209   Homo sapiens EST-YD1 protein (EST-YD1), mRNA   NM   021209   Homo sapiens EST-YD1 protein (EST-YD1), mRNA   NM   021199   Homo sapiens CGI-44 protein; sulfide dehydrogenase like (yeast) (CGI-44), mRNA   NM   021198   Homo sapiens nuclear LIM interactor-interacting factor (NLI-IF), mRNA   NM   021193   Homo sapiens homeo box D12 (HOXD12), mRNA   NM   021193   Homo sapiens homeo box D11 (HOXD11), mRNA   NM   021188   Homo sapiens homeo box D11 (HOXD11), mRNA   NM   021184   Homo sapiens G4 protein (G4), mRNA   NM   021174   Homo sapiens G4 protein (G4), mRNA   NM   021174   Homo sapiens G4 protein (LOC57805), mRNA   NM   021174   Homo sapiens B90 DBC protein (LOC57805), mRNA   NM   021159   Homo sapiens RAP1, GTP-GDP dissociation stimulator 1 (RAP1GDS1), mRNA   NM   021147   Homo sapiens CD209 antigen (CD209), mRNA   NM   021147   Homo sapiens ubiquitously transcribed tetratricopeptide repeat gene, X   chromosome (UTX), mRNA   NM   021139   Homo sapiens UDP glycosyltransferase 2 family, polypeptide B4 (UGT2B4), mRNA   NM   021136   Homo sapiens TNF receptor-associated factor 2 (TRAF2), mRNA   NM   021137   Homo sapiens tumor necrosis factor, alpha-induced protein 1 (endothelial) (TNFAIP1), mRNA   Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA   NM   021130   Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA   NM   021130   Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA   NM   021130   Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA   NM   021130   Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA   Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA   Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA   Homo sapiens discs, larg		Homo sapiens TERA protein (TERA), mRNA
NM 021201 Homo sapiens G5b protein (G5B), mRNA NM 021210 Homo sapiens MUM2 protein (MUM2), mRNA NM 021208 Homo sapiens EST-YD1 protein (EST-YD1), mRNA NM 021200 Homo sapiens PH domain containing protein in retina 1 (PHRET1), mRNA NM 021199 Homo sapiens CGI-44 protein; sulfide dehydrogenase like (yeast) (CGI-44), mRNA NM 021198 Homo sapiens nuclear LIM interactor-interacting factor (NLI-IF), mRNA NM 021193 Homo sapiens homeo box D12 (HOXD12), mRNA NM 021194 Homo sapiens homeo box D11 (HOXD11), mRNA NM 021185 Homo sapiens clones 23667 and 23775 zinc finger protein (LOC57862), mRNA NM 021186 Homo sapiens G4 protein (G4), mRNA NM 021187 Homo sapiens G9 protein (G4), mRNA NM 021188 Homo sapiens BOBC protein (LOC57805), mRNA NM 021170 Homo sapiens p30 DBC protein (LOC57805), mRNA NM 021171 Homo sapiens p30 DBC protein (LOC57805), mRNA NM 021167 Homo sapiens RAP1, GTP-GDP dissociation stimulator 1 (RAP1GDS1), mRNA NM 021159 Homo sapiens CD209 antigen (CD209), mRNA NM 021147 Homo sapiens uracil-DNA glycosylase 2 (UNG2), mRNA NM 021147 Homo sapiens UPP glycosyltransferase 2 family, polypeptide B4 (UGT2B4), mRNA NM 021138 Homo sapiens TNF receptor-associated factor 2 (TRAF2), mRNA NM 021137 Homo sapiens TNF receptor-associated factor 2 (TRAF2), mRNA NM 021136 Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA NM 021131 Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA NM 021130 Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA NM 021130 Homo sapiens sibosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA NM 021130 Homo sapiens sibosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA NM 021130 Homo sapiens sibonuclease L (2',5'-oligoisoadenylate synthetase-dependent) (RNASEL), mRNA NM 02120 Homo sapiens sibosomal protein some receptor interactor 11 (TRIP11), mRNA NM 02120 Homo sapiens siboroid hormone receptor interactor 12 (TRIP12), mRNA NM 004239 Homo sapiens thyroid hormone receptor interactor 12 (TRIP12), mRNA NM 004745		Homo sapiens myosin light chain 2a (LOC58498), mRNA
NM 02120 Homo sapiens MUM2 protein (MUM2), mRNA NM 021200 Homo sapiens EST-YD1 protein (EST-YD1), mRNA NM 021199 Homo sapiens CGI-44 protein; sulfide dehydrogenase like (yeast) (CGI-44), mRNA NM 021198 Homo sapiens nuclear LIM interactor-interacting factor (NLI-IF), mRNA NM 021193 Homo sapiens homeo box D12 (HOXD12), mRNA NM 021194 Homo sapiens homeo box D11 (HOXD11), mRNA NM 021195 Homo sapiens clones 23667 and 23775 zinc finger protein (LOC57862), mRNA NM 021186 Homo sapiens G4 protein (G4), mRNA NM 021177 Homo sapiens U6 snRNA-associated Sm-like protein (LSM2), mRNA NM 021174 Homo sapiens p30 DBC protein (LOC57805), mRNA NM 021175 Homo sapiens p30 DBC protein (LOC57805), mRNA NM 021167 Homo sapiens RAP1, GTP-GDP dissociation stimulator 1 (RAP1GDS1), mRNA NM 021147 Homo sapiens uracil-DNA glycosylase 2 (UNG2), mRNA NM 021140 Homo sapiens uracil-DNA glycosylase 2 (UNG2), mRNA NM 021130 Homo sapiens uracil-DNA glycosylase 2 (UNG2), mRNA NM 021131 Homo sapiens uracil-DNA glycosylase 2 (UNG2), mRNA NM 021130 Homo sapiens uracil-DNA glycosylase 2 (UNG2), mRNA NM 021131 Homo sapiens uracil-DNA glycosylase 2 (UNG2), mRNA NM 021131 Homo sapiens uracil-DNA glycosylase 2 (UNG2), mRNA NM 021131 Homo sapiens uracil-DNA glycosylase 2 (UNG2), mRNA NM 021131 Homo sapiens uracil-DNA glycosylase 2 (UNG2), mRNA NM 021131 Homo sapiens uracil-DNA glycosylase 2 (UNG2), mRNA NM 021131 Homo sapiens uracil-DNA glycosylase 2 (UNG2), mRNA NM 021131 Homo sapiens uracil-DNA glycosylase 2 (UNG2), mRNA NM 021131 Homo sapiens uracil-DNA glycosylase 2 (UNG2), mRNA NM 021131 Homo sapiens uracil-DNA glycosylase 2 (UNG2), mRNA NM 021130 Homo sapiens uracil-DNA glycosylase 2 (UNG2), mRNA NM 021131 Homo sapiens uracil-DNA glycosylase 2 (UNG2), mRNA NM 021130 Homo sapiens through duraced protein 1 (endothelial) (TNFAIP1), mRNA NM 021131 Homo sapiens reticulon 1 (RTN1), mRNA NM 021130 Homo sapiens reticulon 1 (RTN1), mRNA NM 021131 Homo sapiens reticulon 1 (RTN1), mRNA NM 021130 Homo sapiens peptidylprolyl isomerase A (cyclophilin A) (PPIA), mRNA		Homo sapiens G5b protein (G5B), mRNA
NM 021108		Homo sapiens MUM2 protein (MUM2), mRNA
NM 021199   Homo sapiens PH domain containing protein in retina 1 (PHRE11), mRNA   Homo sapiens CGI-44 protein; sulfide dehydrogenase like (yeast) (CGI-44), mRNA   NM 021193   Homo sapiens nuclear LIM interactor-interacting factor (NLI-IF), mRNA   NM 021194   Homo sapiens homeo box D12 (HOXD12), mRNA   NM 021188   Homo sapiens homeo box D11 (HOXD11), mRNA   NM 021184   Homo sapiens Cd protein (G4), mRNA   NM 021174   Homo sapiens U6 snRNA-associated Sm-like protein (LSM2), mRNA   NM 021174   Homo sapiens DBC protein (LOC57805), mRNA   NM 021175   Homo sapiens hypothetical protein WUGSC:H_RG083M05.2 (LOC57798), mRNA   NM 021159   Homo sapiens RAP1, GTP-GDP dissociation stimulator 1 (RAP1GDS1), mRNA   NM 021147   Homo sapiens by postitute of the companies of th		Homo sapiens EST-YD1 protein (EST-YD1), mRNA
NM   021199		Homo sapiens PH domain containing protein in retina 1 (PHRET1), mRNA
mRNA NM 021198 Homo sapiens nuclear LIM interactor-interacting factor (NLI-IF), mRNA NM 021192 Homo sapiens homeo box D12 (HOXD12), mRNA NM 021192 Homo sapiens homeo box D11 (HOXD11), mRNA NM 021188 Homo sapiens clones 23667 and 23775 zinc finger protein (LOC57862), mRNA NM 021184 Homo sapiens U6 snRNA-associated Sm-like protein (LSM2), mRNA NM 021177 Homo sapiens p30 DBC protein (LOC57805), mRNA NM 021174 Homo sapiens p30 DBC protein (LOC57805), mRNA NM 021167 Homo sapiens p30 DBC protein (LOC57805), mRNA NM 021167 Homo sapiens RAP1, GTP-GDP dissociation stimulator 1 (RAP1GDS1), mRNA NM 021159 Homo sapiens CD209 antigen (CD209), mRNA NM 021147 Homo sapiens uracil-DNA glycosylase 2 (UNG2), mRNA NM 021140 Homo sapiens uracil-DNA glycosylase 2 (UNG2), mRNA NM_021139 Homo sapiens uracil-DNA glycosylase 2 (UNG2), mRNA NM_021139 Homo sapiens UDP glycosyltransferase 2 family, polypeptide B4 (UGT2B4), mRNA NM_021138 Homo sapiens TNF receptor-associated factor 2 (TRAF2), mRNA NM_021137 Homo sapiens tumor necrosis factor, alpha-induced protein 1 (endothelial) (TNFAIP1), mRNA NM_021135 Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA NM_021135 Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA NM_021130 Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA NM_021130 Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA NM_021130 Homo sapiens peptidylprolyl isomerase A (cyclophilin A) (PPIA), mRNA NM_021130 Homo sapiens peptidylprolyl isomerase A (cyclophilin A) (PPIA), mRNA NM_021130 Homo sapiens sthyroid hormone receptor interactor 11 (TRIP11), mRNA NM_04239 Homo sapiens thyroid hormone receptor interactor 12 (TRIP12), mRNA NM_04238 Homo sapiens thyroid hormone receptor interactor 12 (TRIP12), mRNA NM_04454 Homo sapiens discs, large (Drosophila) homolog-associated protein 2		Homo sapiens CGI-44 protein: sulfide dehydrogenase like (yeast) (CGI-44),
NM 021198 Homo sapiens nuclear LIM interactor-interacting factor (NLI-IF), mRNA NM 021193 Homo sapiens homeo box D12 (HOXD12), mRNA NM 021198 Homo sapiens homeo box D11 (HOXD11), mRNA NM 021188 Homo sapiens clones 23667 and 23775 zinc finger protein (LOC57862), mRNA NM 021187 Homo sapiens G4 protein (G4), mRNA NM 021177 Homo sapiens U6 snRNA-associated Sm-like protein (LSM2), mRNA NM 021167 Homo sapiens p30 DBC protein (LOC57805), mRNA NM 021167 Homo sapiens hypothetical protein WUGSC:H_RG083M05.2 (LOC57798), mRNA NM 021159 Homo sapiens RAP1, GTP-GDP dissociation stimulator 1 (RAP1GDS1), mRNA NM 021140 Homo sapiens uracil-DNA glycosylase 2 (UNG2), mRNA NM 021140 Homo sapiens ubiquitously transcribed tetratricopeptide repeat gene, X chromosome (UTX), mRNA NM 021139 Homo sapiens UDP glycosyltransferase 2 family, polypeptide B4 (UGT2B4), mRNA NM 021138 Homo sapiens TNF receptor-associated factor 2 (TRAF2), mRNA NM 021137 Homo sapiens tumor necrosis factor, alpha-induced protein 1 (endothelial) (TNFAIP1), mRNA NM 021135 Homo sapiens reticulon 1 (RTN1), mRNA NM 021136 Homo sapiens reticulon 1 (RTN1), mRNA NM 021137 Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA NM 021130 Homo sapiens ribonuclease L (2',5'-oligoisoadenylate synthetase-dependent) (RNASEL), mRNA NM 021130 Homo sapiens peptidylprolyl isomerase A (cyclophilin A) (PPIA), mRNA NM 021130 Homo sapiens sthyroid hormone receptor interactor 11 (TRIP11), mRNA NM 004239 Homo sapiens thyroid hormone receptor interactor 12 (TRIP12), mRNA NM 004238 Homo sapiens thyroid hormone receptor interactor 12 (TRIP12), mRNA NM 004745 Homo sapiens discs, large (Drosophila) homolog-associated protein 2	14141_021199	mRNA
NM 021193 Homo sapiens homeo box D12 (HOXD12), mRNA NM 021188 Homo sapiens homeo box D11 (HOXD11), mRNA NM 021188 Homo sapiens clones 23667 and 23775 zinc finger protein (LOC57862), mRNA NM 021184 Homo sapiens G4 protein (G4), mRNA NM 021177 Homo sapiens U6 snRNA-associated Sm-like protein (LSM2), mRNA NM 021174 Homo sapiens p30 DBC protein (LOC57805), mRNA NM 021167 Homo sapiens hypothetical protein WUGSC:H_RG083M05.2 (LOC57798), mRNA NM 021159 Homo sapiens RAP1, GTP-GDP dissociation stimulator 1 (RAP1GDS1), mRNA NM 021155 Homo sapiens CD209 antigen (CD209), mRNA NM 021147 Homo sapiens wacil-DNA glycosylase 2 (UNG2), mRNA NM 021140 Homo sapiens unacil-DNA glycosylase 2 (UNG2), mRNA NM 021140 Homo sapiens UDP glycosyltranscribed tetratricopeptide repeat gene, X chromosome (UTX), mRNA NM 021138 Homo sapiens TNF receptor-associated factor 2 (TRAF2), mRNA NM 021136 Homo sapiens tumor necrosis factor, alpha-induced protein 1 (endothelial) (TNFAIP1), mRNA NM 021135 Homo sapiens reticulon 1 (RTN1), mRNA NM 021136 Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA NM 021130 Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA NM 021130 Homo sapiens geptidylprolyl isomerase A (cyclophilin A) (PPIA), mRNA NM 021130 Homo sapiens discs, large (Drosophila) homolog 3 (neuroendocrine-dlg) (DLG3), mRNA NM 004239 Homo sapiens thyroid hormone receptor interactor 11 (TRIP11), mRNA NM 004238 Homo sapiens discs, large (Drosophila) homolog-associated protein 2	NM 021198	Homo sapiens nuclear LIM interactor-interacting factor (NLI-IF), mRNA
NM 021182 Homo sapiens homeo box D11 (HOXD11), mRNA  NM 021188 Homo sapiens clones 23667 and 23775 zinc finger protein (LOC57862), mRNA  NM 021177 Homo sapiens U6 snRNA-associated Sm-like protein (LSM2), mRNA  NM 021174 Homo sapiens p30 DBC protein (LOC57805), mRNA  NM 021167 Homo sapiens hypothetical protein WUGSC:H_RG083M05.2 (LOC57798), mRNA  NM 021159 Homo sapiens RAP1, GTP-GDP dissociation stimulator 1 (RAP1GDS1), mRNA  NM 021155 Homo sapiens wracil-DNA glycosylase 2 (UNG2), mRNA  NM 021147 Homo sapiens ubiquitously transcribed tetratricopeptide repeat gene, X chromosome (UTX), mRNA  NM 021139 Homo sapiens UDP glycosyltransferase 2 family, polypeptide B4 (UGT2B4), mRNA  NM 021137 Homo sapiens tumor necrosis factor, alpha-induced protein 1 (endothelial) (TNFAIP1), mRNA  NM 021136 Homo sapiens reticulon 1 (RTN1), mRNA  NM 021137 Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA  NM 021130 Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA  NM 021130 Homo sapiens peptidylprolyl isomerase A (cyclophilin A) (PPIA), mRNA  NM 021130 Homo sapiens peptidylprolyl isomerase A (cyclophilin A) (PPIA), mRNA  NM 021130 Homo sapiens peptidylprolyl isomerase A (cyclophilin A) (PPIA), mRNA  NM 021130 Homo sapiens peptidylprolyl isomerase A (cyclophilin A) (PPIA), mRNA  NM 02120 Homo sapiens thyroid hormone receptor interactor 11 (TRIP11), mRNA  NM 004239 Homo sapiens thyroid hormone receptor interactor 12 (TRIP12), mRNA  NM 004238 Homo sapiens discs, large (Drosophila) homolog-associated protein 2		Homo sapiens homeo box D12 (HOXD12), mRNA
NM 021188 Homo sapiens clones 23667 and 23775 zinc finger protein (LOC57862), mRNA NM 021174 Homo sapiens G4 protein (G4), mRNA NM 021174 Homo sapiens U6 snRNA-associated Sm-like protein (LSM2), mRNA NM 021167 Homo sapiens p30 DBC protein (LOC57805), mRNA NM 021167 Homo sapiens hypothetical protein WUGSC:H_RG083M05.2 (LOC57798), mRNA NM 021159 Homo sapiens RAP1, GTP-GDP dissociation stimulator 1 (RAP1GDS1), mRNA NM 021155 Homo sapiens UD209 antigen (CD209), mRNA NM 021147 Homo sapiens uracil-DNA glycosylase 2 (UNG2), mRNA NM 021140 Homo sapiens ubiquitously transcribed tetratricopeptide repeat gene, X chromosome (UTX), mRNA NM 021139 Homo sapiens UDP glycosyltransferase 2 family, polypeptide B4 (UGT2B4), mRNA NM 021136 Homo sapiens TNF receptor-associated factor 2 (TRAF2), mRNA NM 021137 Homo sapiens tumor necrosis factor, alpha-induced protein 1 (endothelial) (TNFAIP1), mRNA NM 021135 Homo sapiens reticulon 1 (RTN1), mRNA NM 021136 Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA NM 021130 Homo sapiens ribonuclease L (2',5'-oligoisoadenylate synthetase-dependent) (RNASEL), mRNA NM 021120 Homo sapiens peptidylprolyl isomerase A (cyclophilin A) (PPIA), mRNA NM 02120 Homo sapiens discs, large (Drosophila) homolog 3 (neuroendocrine-dlg) (DLG3), mRNA NM 004238 Homo sapiens thyroid hormone receptor interactor 11 (TRIP11), mRNA NM 004238 Homo sapiens thyroid hormone receptor interactor 12 (TRIP12), mRNA NM 004745 Homo sapiens discs, large (Drosophila) homolog-associated protein 2		Homo sapiens homeo box D11 (HOXD11), mRNA
NM_021174 Homo sapiens G4 protein (G4), mRNA NM_021174 Homo sapiens U6 snRNA-associated Sm-like protein (LSM2), mRNA NM_021167 Homo sapiens p30 DBC protein (LOC57805), mRNA NM_021167 Homo sapiens hypothetical protein WUGSC:H_RG083M05.2 (LOC57798), mRNA NM_021159 Homo sapiens RAP1, GTP-GDP dissociation stimulator 1 (RAP1GDS1), mRNA NM_021155 Homo sapiens UD209 antigen (CD209), mRNA NM_021140 Homo sapiens uracil-DNA glycosylase 2 (UNG2), mRNA NM_021140 Homo sapiens ubiquitously transcribed tetratricopeptide repeat gene, X chromosome (UTX), mRNA NM_021139 Homo sapiens UDP glycosyltransferase 2 family, polypeptide B4 (UGT2B4), mRNA NM_021136 Homo sapiens TNF receptor-associated factor 2 (TRAF2), mRNA NM_021136 Homo sapiens reticulon 1 (RTN1), mRNA NM_021136 Homo sapiens reticulon 1 (RTN1), mRNA NM_021137 Homo sapiens reticulon 1 (RTN1), mRNA NM_021138 Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA NM_021130 Homo sapiens ribonuclease L (2',5'-oligoisoadenylate synthetase-dependent) (RNASEL), mRNA NM_021130 Homo sapiens peptidylprolyl isomerase A (cyclophilin A) (PPIA), mRNA NM_021120 Homo sapiens discs, large (Drosophila) homolog 3 (neuroendocrine-dlg) (DLG3), mRNA NM_004238 Homo sapiens thyroid hormone receptor interactor 11 (TRIP11), mRNA NM_004238 Homo sapiens thyroid hormone receptor interactor 12 (TRIP12), mRNA NM_004745 Homo sapiens discs, large (Drosophila) homolog-associated protein 2		Homo sapiens clones 23667 and 23775 zinc finger protein (LOC57862), mRNA
NM 021177 Homo sapiens U6 snRNA-associated Sm-like protein (LSM2), mRNA NM 021167 Homo sapiens p30 DBC protein (LOC57805), mRNA NM 021167 Homo sapiens hypothetical protein WUGSC:H_RG083M05.2 (LOC57798), mRNA NM 021159 Homo sapiens RAP1, GTP-GDP dissociation stimulator 1 (RAP1GDS1), mRNA NM 021155 Homo sapiens CD209 antigen (CD209), mRNA NM 021147 Homo sapiens uracil-DNA glycosylase 2 (UNG2), mRNA NM 021140 Homo sapiens ubiquitously transcribed tetratricopeptide repeat gene, X chromosome (UTX), mRNA NM 021139 Homo sapiens UDP glycosyltransferase 2 family, polypeptide B4 (UGT2B4), mRNA NM 021138 Homo sapiens TNF receptor-associated factor 2 (TRAF2), mRNA NM 021136 Homo sapiens ribosomal protein 36 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA NM 021131 Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA NM 021130 Homo sapiens ribonuclease L (2',5'-oligoisoadenylate synthetase-dependent) (RNASEL), mRNA NM 021130 Homo sapiens peptidylprolyl isomerase A (cyclophilin A) (PPIA), mRNA NM 021120 Homo sapiens discs, large (Drosophila) homolog 3 (neuroendocrine-dlg) (DLG3), mRNA NM 004238 Homo sapiens thyroid hormone receptor interactor 11 (TRIP11), mRNA NM 004238 Homo sapiens discs, large (Drosophila) homolog-associated protein 2		Homo sapiens G4 protein (G4), mRNA
NM_021174 Homo sapiens p30 DBC protein (LOC57805), mRNA  NM_021167 Homo sapiens hypothetical protein WUGSC:H_RG083M05.2 (LOC57798), mRNA  NM_021159 Homo sapiens RAP1, GTP-GDP dissociation stimulator 1 (RAP1GDS1), mRNA  NM_021155 Homo sapiens CD209 antigen (CD209), mRNA  NM_021147 Homo sapiens uracil-DNA glycosylase 2 (UNG2), mRNA  NM_021140 Homo sapiens ubiquitously transcribed tetratricopeptide repeat gene, X chromosome (UTX), mRNA  NM_021139 Homo sapiens UDP glycosyltransferase 2 family, polypeptide B4 (UGT2B4), mRNA  NM_021137 Homo sapiens tumor necrosis factor, alpha-induced protein 1 (endothelial) (TNFAIP1), mRNA  NM_021136 Homo sapiens reticulon 1 (RTN1), mRNA  NM_021135 Homo sapiens reticulon 1 (RTN1), mRNA  NM_021136 Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA  NM_021130 Homo sapiens ribonuclease L (2',5'-oligoisoadenylate synthetase-dependent) (RNASEL), mRNA  NM_021130 Homo sapiens peptidylprolyl isomerase A (cyclophilin A) (PPIA), mRNA  NM_021120 Homo sapiens discs, large (Drosophila) homolog 3 (neuroendocrine-dlg) (DLG3), mRNA  NM_004238 Homo sapiens thyroid hormone receptor interactor 11 (TRIP11), mRNA  NM_004238 Homo sapiens thyroid hormone receptor interactor 12 (TRIP12), mRNA  NM_004745 Homo sapiens discs, large (Drosophila) homolog-associated protein 2		Homo sapiens LI6 snRNA-associated Sm-like protein (LSM2), mRNA
NM_021157 Homo sapiens hypothetical protein WUGSC:H_RG083M05.2 (LOC57798), mRNA  NM_021159 Homo sapiens RAP1, GTP-GDP dissociation stimulator 1 (RAP1GDS1), mRNA  NM_021147 Homo sapiens uracil-DNA glycosylase 2 (UNG2), mRNA  NM_021140 Homo sapiens ubiquitously transcribed tetratricopeptide repeat gene, X chromosome (UTX), mRNA  NM_021139 Homo sapiens UDP glycosyltransferase 2 family, polypeptide B4 (UGT2B4), mRNA  NM_021138 Homo sapiens TNF receptor-associated factor 2 (TRAF2), mRNA  NM_021137 Homo sapiens tumor necrosis factor, alpha-induced protein 1 (endothelial) (TNFAIP1), mRNA  NM_021136 Homo sapiens reticulon 1 (RTN1), mRNA  NM_021135 Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA  NM_021130 Homo sapiens ribonuclease L (2',5'-oligoisoadenylate synthetase-dependent) (RNASEL), mRNA  NM_021130 Homo sapiens geptidylprolyl isomerase A (cyclophilin A) (PPIA), mRNA  NM_021120 Homo sapiens discs, large (Drosophila) homolog 3 (neuroendocrine-dlg) (DLG3), mRNA  NM_004239 Homo sapiens thyroid hormone receptor interactor 11 (TRIP11), mRNA  NM_004238 Homo sapiens discs, large (Drosophila) homolog-associated protein 2		Home sapiens n30 DBC protein (LOC57805), mRNA
NM 021159 Homo sapiens RAP1, GTP-GDP dissociation stimulator 1 (RAP1GDS1), mRNA NM 021147 Homo sapiens uracil-DNA glycosylase 2 (UNG2), mRNA NM 021140 Homo sapiens ubiquitously transcribed tetratricopeptide repeat gene, X chromosome (UTX), mRNA NM 021139 Homo sapiens UDP glycosyltransferase 2 family, polypeptide B4 (UGT2B4), mRNA NM 021138 Homo sapiens TNF receptor-associated factor 2 (TRAF2), mRNA NM 021137 Homo sapiens tumor necrosis factor, alpha-induced protein 1 (endothelial) (TNFAIP1), mRNA NM 021136 Homo sapiens reticulon 1 (RTN1), mRNA NM 021135 Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA NM 021130 Homo sapiens ribonuclease L (2',5'-oligoisoadenylate synthetase-dependent) (RNASEL), mRNA NM 021130 Homo sapiens peptidylprolyl isomerase A (cyclophilin A) (PPIA), mRNA NM 021120 Homo sapiens discs, large (Drosophila) homolog 3 (neuroendocrine-dlg) (DLG3), mRNA NM 004239 Homo sapiens thyroid hormone receptor interactor 11 (TRIP11), mRNA NM 004238 Homo sapiens thyroid hormone receptor interactor 12 (TRIP12), mRNA NM 004745 Homo sapiens discs, large (Drosophila) homolog-associated protein 2		Home sapiens by other ical protein WUGSC:H RG083M05.2 (LOC57798),
NM 021159 Homo sapiens RAP1, GTP-GDP dissociation stimulator 1 (RAP1GDS1), mRNA  NM 021147 Homo sapiens UD209 antigen (CD209), mRNA  NM 021140 Homo sapiens uracil-DNA glycosylase 2 (UNG2), mRNA  NM 021130 Homo sapiens ubiquitously transcribed tetratricopeptide repeat gene, X chromosome (UTX), mRNA  NM 021139 Homo sapiens UDP glycosyltransferase 2 family, polypeptide B4 (UGT2B4), mRNA  NM 021138 Homo sapiens TNF receptor-associated factor 2 (TRAF2), mRNA  NM 021137 Homo sapiens tumor necrosis factor, alpha-induced protein 1 (endothelial) (TNFAIP1), mRNA  NM 021135 Homo sapiens reticulon 1 (RTN1), mRNA  NM 021135 Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA  NM 021130 Homo sapiens ribonuclease L (2',5'-oligoisoadenylate synthetase-dependent) (RNASEL), mRNA  NM 021120 Homo sapiens discs, large (Drosophila) homolog 3 (neuroendocrine-dlg) (DLG3), mRNA  NM 004239 Homo sapiens thyroid hormone receptor interactor 11 (TRIP11), mRNA  NM 004238 Homo sapiens discs, large (Drosophila) homolog-associated protein 2	NWI_021107	mRNA
NM_021147 Homo sapiens CD209 antigen (CD209), mRNA NM_021140 Homo sapiens uracil-DNA glycosylase 2 (UNG2), mRNA NM_021130 Homo sapiens ubiquitously transcribed tetratricopeptide repeat gene, X chromosome (UTX), mRNA NM_021139 Homo sapiens UDP glycosyltransferase 2 family, polypeptide B4 (UGT2B4), mRNA NM_021138 Homo sapiens TNF receptor-associated factor 2 (TRAF2), mRNA NM_021137 Homo sapiens tumor necrosis factor, alpha-induced protein 1 (endothelial) (TNFAIP1), mRNA NM_021136 Homo sapiens reticulon 1 (RTN1), mRNA NM_021135 Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA NM_021131 Homo sapiens ribonuclease L (2',5'-oligoisoadenylate synthetase-dependent) (RNASEL), mRNA NM_021120 Homo sapiens peptidylprolyl isomerase A (cyclophilin A) (PPIA), mRNA NM_021120 Homo sapiens discs, large (Drosophila) homolog 3 (neuroendocrine-dlg) (DLG3), mRNA NM_004239 Homo sapiens thyroid hormone receptor interactor 11 (TRIP11), mRNA NM_004238 Homo sapiens discs, large (Drosophila) homolog-associated protein 2	NM 021159	Homo sapiens RAP1, GTP-GDP dissociation stimulator 1 (RAP1GDS1), mRNA
NM 021147 Homo sapiens uracil-DNA glycosylase 2 (UNG2), mRNA  NM 021140 Homo sapiens ubiquitously transcribed tetratricopeptide repeat gene, X chromosome (UTX), mRNA  NM 021139 Homo sapiens UDP glycosyltransferase 2 family, polypeptide B4 (UGT2B4), mRNA  NM 021138 Homo sapiens TNF receptor-associated factor 2 (TRAF2), mRNA  NM 021137 Homo sapiens tumor necrosis factor, alpha-induced protein 1 (endothelial) (TNFAIP1), mRNA  NM 021136 Homo sapiens reticulon 1 (RTN1), mRNA  NM 021135 Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA  NM 021130 Homo sapiens ribonuclease L (2',5'-oligoisoadenylate synthetase-dependent) (RNASEL), mRNA  NM 021120 Homo sapiens discs, large (Drosophila) homolog 3 (neuroendocrine-dlg) (DLG3), mRNA  NM 004239 Homo sapiens thyroid hormone receptor interactor 11 (TRIP11), mRNA  NM 004238 Homo sapiens discs, large (Drosophila) homolog-associated protein 2		Homo sapiens CD209 antigen (CD209), mRNA
NM_021130 Homo sapiens ubiquitously transcribed tetratricopeptide repeat gene, X chromosome (UTX), mRNA  NM_021139 Homo sapiens UDP glycosyltransferase 2 family, polypeptide B4 (UGT2B4), mRNA  NM_021138 Homo sapiens TNF receptor-associated factor 2 (TRAF2), mRNA  NM_021137 Homo sapiens tumor necrosis factor, alpha-induced protein 1 (endothelial) (TNFAIP1), mRNA  NM_021136 Homo sapiens reticulon 1 (RTN1), mRNA  NM_021135 Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA  NM_021131 Homo sapiens ribonuclease L (2',5'-oligoisoadenylate synthetase-dependent) (RNASEL), mRNA  NM_021130 Homo sapiens peptidylprolyl isomerase A (cyclophilin A) (PPIA), mRNA  NM_021120 Homo sapiens discs, large (Drosophila) homolog 3 (neuroendocrine-dlg) (DLG3), mRNA  NM_004239 Homo sapiens thyroid hormone receptor interactor 11 (TRIP11), mRNA  NM_004238 Homo sapiens thyroid hormone receptor interactor 12 (TRIP12), mRNA  NM_004745 Homo sapiens discs, large (Drosophila) homolog-associated protein 2		Homo sapiens uracil-DNA glycosylase 2 (UNG2), mRNA
chromosome (UTX), mRNA  NM_021139 Homo sapiens UDP glycosyltransferase 2 family, polypeptide B4 (UGT2B4), mRNA  NM_021138 Homo sapiens TNF receptor-associated factor 2 (TRAF2), mRNA  NM_021137 Homo sapiens tumor necrosis factor, alpha-induced protein 1 (endothelial) (TNFAIP1), mRNA  NM_021136 Homo sapiens reticulon 1 (RTN1), mRNA  NM_021135 Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA  NM_021133 Homo sapiens ribonuclease L (2',5'-oligoisoadenylate synthetase-dependent) (RNASEL), mRNA  NM_021130 Homo sapiens peptidylprolyl isomerase A (cyclophilin A) (PPIA), mRNA  NM_021120 Homo sapiens discs, large (Drosophila) homolog 3 (neuroendocrine-dlg) (DLG3), mRNA  NM_004239 Homo sapiens thyroid hormone receptor interactor 11 (TRIP11), mRNA  NM_004238 Homo sapiens discs, large (Drosophila) homolog-associated protein 2		Homo sapiens ubiquitously transcribed tetratricopeptide repeat gene, X
NM_021139 Homo sapiens UDP glycosyltransferase 2 family, polypeptide B4 (UGT2B4), mRNA  NM_021138 Homo sapiens TNF receptor-associated factor 2 (TRAF2), mRNA  NM_021137 Homo sapiens tumor necrosis factor, alpha-induced protein 1 (endothelial) (TNFAIP1), mRNA  NM_021136 Homo sapiens reticulon 1 (RTN1), mRNA  NM_021135 Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA  NM_021133 Homo sapiens ribonuclease L (2',5'-oligoisoadenylate synthetase-dependent) (RNASEL), mRNA  NM_021130 Homo sapiens peptidylprolyl isomerase A (cyclophilin A) (PPIA), mRNA  NM_021120 Homo sapiens discs, large (Drosophila) homolog 3 (neuroendocrine-dlg) (DLG3), mRNA  NM_004239 Homo sapiens thyroid hormone receptor interactor 11 (TRIP11), mRNA  NM_004238 Homo sapiens thyroid hormone receptor interactor 12 (TRIP12), mRNA  NM_004745 Homo sapiens discs, large (Drosophila) homolog-associated protein 2	14141_021140	chromosome (UTX) mRNA
mRNA  NM 021138 Homo sapiens TNF receptor-associated factor 2 (TRAF2), mRNA  NM_021137 Homo sapiens tumor necrosis factor, alpha-induced protein 1 (endothelial)  (TNFAIP1), mRNA  NM_021136 Homo sapiens reticulon 1 (RTN1), mRNA  NM_021135 Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA  NM_021133 Homo sapiens ribonuclease L (2',5'-oligoisoadenylate synthetase-dependent)  (RNASEL), mRNA  NM_021130 Homo sapiens peptidylprolyl isomerase A (cyclophilin A) (PPIA), mRNA  NM_021120 Homo sapiens discs, large (Drosophila) homolog 3 (neuroendocrine-dlg)  (DLG3), mRNA  NM_004239 Homo sapiens thyroid hormone receptor interactor 11 (TRIP11), mRNA  NM_004238 Homo sapiens discs, large (Drosophila) homolog-associated protein 2	NM_021139	Homo sapiens UDP glycosyltransferase 2 family, polypeptide B4 (UGT2B4),
NM_021137 Homo sapiens tumor necrosis factor, alpha-induced protein 1 (endothelial)  (TNFAIP1), mRNA  NM_021136 Homo sapiens reticulon 1 (RTN1), mRNA  NM_021135 Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA  NM_021133 Homo sapiens ribonuclease L (2',5'-oligoisoadenylate synthetase-dependent)  (RNASEL), mRNA  NM_021130 Homo sapiens peptidylprolyl isomerase A (cyclophilin A) (PPIA), mRNA  NM_021120 Homo sapiens discs, large (Drosophila) homolog 3 (neuroendocrine-dlg)  (DLG3), mRNA  NM_004239 Homo sapiens thyroid hormone receptor interactor 11 (TRIP11), mRNA  NM_004238 Homo sapiens thyroid hormone receptor interactor 12 (TRIP12), mRNA  NM_004745 Homo sapiens discs, large (Drosophila) homolog-associated protein 2		mRNA PNA
(TNFAIP1), mRNA  NM_021136 Homo sapiens reticulon 1 (RTN1), mRNA  NM_021135 Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA  NM_021133 Homo sapiens ribonuclease L (2',5'-oligoisoadenylate synthetase-dependent) (RNASEL), mRNA  NM_021130 Homo sapiens peptidylprolyl isomerase A (cyclophilin A) (PPIA), mRNA  NM_021120 Homo sapiens discs, large (Drosophila) homolog 3 (neuroendocrine-dlg) (DLG3), mRNA  NM_004239 Homo sapiens thyroid hormone receptor interactor 11 (TRIP11), mRNA  NM_004238 Homo sapiens thyroid hormone receptor interactor 12 (TRIP12), mRNA  NM_004745 Homo sapiens discs, large (Drosophila) homolog-associated protein 2	NM_021138	Homo sapiens TNF receptor-associated factor 2 (TRAF2), mRNA
NM_021136 Homo sapiens reticulon 1 (RTN1), mRNA  NM_021135 Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA  NM_021133 Homo sapiens ribonuclease L (2',5'-oligoisoadenylate synthetase-dependent) (RNASEL), mRNA  NM_021130 Homo sapiens peptidylprolyl isomerase A (cyclophilin A) (PPIA), mRNA  NM_021120 Homo sapiens discs, large (Drosophila) homolog 3 (neuroendocrine-dlg) (DLG3), mRNA  NM_004239 Homo sapiens thyroid hormone receptor interactor 11 (TRIP11), mRNA  NM_004238 Homo sapiens thyroid hormone receptor interactor 12 (TRIP12), mRNA  NM_004745 Homo sapiens discs, large (Drosophila) homolog-associated protein 2	NM_021137	Homo sapiens tumor necrosis factor, alpha-induced protein 1 (endothelial) (TNFAIP1), mRNA
NM_021135 Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2), mRNA  NM_021133 Homo sapiens ribonuclease L (2',5'-oligoisoadenylate synthetase-dependent) (RNASEL), mRNA  NM_021130 Homo sapiens peptidylprolyl isomerase A (cyclophilin A) (PPIA), mRNA  NM_021120 Homo sapiens discs, large (Drosophila) homolog 3 (neuroendocrine-dlg) (DLG3), mRNA  NM_004239 Homo sapiens thyroid hormone receptor interactor 11 (TRIP11), mRNA  NM_004238 Homo sapiens thyroid hormone receptor interactor 12 (TRIP12), mRNA  NM_004745 Homo sapiens discs, large (Drosophila) homolog-associated protein 2	NM 021136	Homo saniens reticulon 1 (RTN1), mRNA
mRNA  NM_021133 Homo sapiens ribonuclease L (2',5'-oligoisoadenylate synthetase-dependent)  (RNASEL), mRNA  NM_021130 Homo sapiens peptidylprolyl isomerase A (cyclophilin A) (PPIA), mRNA  NM_021120 Homo sapiens discs, large (Drosophila) homolog 3 (neuroendocrine-dlg)  (DLG3), mRNA  NM_004239 Homo sapiens thyroid hormone receptor interactor 11 (TRIP11), mRNA  NM_004238 Homo sapiens thyroid hormone receptor interactor 12 (TRIP12), mRNA  NM_004745 Homo sapiens discs, large (Drosophila) homolog-associated protein 2		Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2),
NM_021133 Homo sapiens ribonuclease L (2',5'-oligoisoadenylate synthetase-dependent) (RNASEL), mRNA  NM_021130 Homo sapiens peptidylprolyl isomerase A (cyclophilin A) (PPIA), mRNA  NM_021120 Homo sapiens discs, large (Drosophila) homolog 3 (neuroendocrine-dlg) (DLG3), mRNA  NM_004239 Homo sapiens thyroid hormone receptor interactor 11 (TRIP11), mRNA  NM_004238 Homo sapiens thyroid hormone receptor interactor 12 (TRIP12), mRNA  NM_004745 Homo sapiens discs, large (Drosophila) homolog-associated protein 2	14141_021133	mRNA
(RNASEL), mRNA  NM 021130 Homo sapiens peptidylprolyl isomerase A (cyclophilin A) (PPIA), mRNA  NM 021120 Homo sapiens discs, large (Drosophila) homolog 3 (neuroendocrine-dlg) (DLG3), mRNA  NM 004239 Homo sapiens thyroid hormone receptor interactor 11 (TRIP11), mRNA  NM 004238 Homo sapiens thyroid hormone receptor interactor 12 (TRIP12), mRNA  NM 004745 Homo sapiens discs, large (Drosophila) homolog-associated protein 2	NM 021133	Homo sapiens ribonuclease L (2',5'-oligoisoadenylate synthetase-dependent)
NM_021130 Homo sapiens peptidylprolyl isomerase A (cyclophilin A) (PPIA), mRNA  NM_021120 Homo sapiens discs, large (Drosophila) homolog 3 (neuroendocrine-dlg) (DLG3), mRNA  NM_004239 Homo sapiens thyroid hormone receptor interactor 11 (TRIP11), mRNA  NM_004238 Homo sapiens thyroid hormone receptor interactor 12 (TRIP12), mRNA  NM_004745 Homo sapiens discs, large (Drosophila) homolog-associated protein 2	11111_021100	(RNASEL), mRNA
NM 021120 Homo sapiens discs, large (Drosophila) homolog 3 (neuroendocrine-dlg) (DLG3), mRNA  NM 004239 Homo sapiens thyroid hormone receptor interactor 11 (TRIP11), mRNA  NM 004238 Homo sapiens thyroid hormone receptor interactor 12 (TRIP12), mRNA  NM 004745 Homo sapiens discs, large (Drosophila) homolog-associated protein 2	NM 021130	Homo sapiens peptidylprolyl isomerase A (cyclophilin A) (PPIA), mRNA
(DLG3), mRNA  NM_004239 Homo sapiens thyroid hormone receptor interactor 11 (TRIP11), mRNA  NM_004238 Homo sapiens thyroid hormone receptor interactor 12 (TRIP12), mRNA  NM_004745 Homo sapiens discs, large (Drosophila) homolog-associated protein 2		Homo sapiens discs, large (Drosophila) homolog 3 (neuroendocrine-dlg)
NM 004238 Homo sapiens thyroid hormone receptor interactor 12 (TRIP12), mRNA NM 004745 Homo sapiens discs, large (Drosophila) homolog-associated protein 2		(DLG3), mRNA
NM 004238 Homo sapiens thyroid hormone receptor interactor 12 (TRIP12), mRNA NM 004745 Homo sapiens discs, large (Drosophila) homolog-associated protein 2	NM_004239	Homo sapiens thyroid hormone receptor interactor 11 (TRIP11), mRNA
NM 004745 Homo sapiens discs, large (Drosophila) homolog-associated protein 2	NM 004238	Homo sapiens thyroid hormone receptor interactor 12 (TRIP12), mRNA
		Homo sapiens discs, large (Drosophila) homolog-associated protein 2
NM_004687 Homo sapiens myotubularin related protein 4 (MTMR4), mRNA	NIM 004687	Homo sapiens myotubularin related protein 4 (MTMR4), mRNA

NM_02105	NM 004348	Homo sapiens runt-related transcription factor 2 (RUNX2), mRNA
(CACNA11), mRNA  NM 021105  Homo sapiens phospholipid scramblase 1 (PLSCR1), mRNA  NM 002957  Homo sapiens requiem, apoptosis response zinc finger gene (REQ), mRNA  NM 001106  Homo sapiens activin A receptor, type II (ACVR2), mRNA  NM 001105  Homo sapiens activin A receptor, type II (ACVR2), mRNA  NM 001105  Homo sapiens activin A receptor, type II (ACVR2), mRNA  NM 001570  Homo sapiens lectin, mannose-binding, 1 (LMAN1), mRNA  NM 013283  Homo sapiens sell blood group precursor (McLeod phenotype) (XK), mRNA  NM 013284  Homo sapiens sell blood group precursor (McLeod phenotype) (XK), mRNA  NM 013258  Homo sapiens small proline-rich protein 2C (SPRR2C), mRNA  NM 006518  Homo sapiens regenerating islet-derived 1 beta (pancreatic stone protein, pancreatic thread protein) (REG1B), mRNA  NM 006534  Homo sapiens Kruppel-like factor 1 (erythroid) (KLF1), mRNA  NM 006353  Homo sapiens Kruppel-like factor 1 (erythroid) (KLF1), mRNA  NM 006354  Homo sapiens mall proline-rich protein 1A (SPRR1A), mRNA  NM 005950  Homo sapiens small proline-rich protein 1A (SPRR1A), mRNA  NM 005951  Homo sapiens small proline-rich protein 1A (SPRR1A), mRNA  NM 005952  Homo sapiens small proline-rich protein (IL18BP), mRNA  NM 005950  Homo sapiens metallothionein 1X (MT1X), mRNA  NM 006618  Homo sapiens metallothionein 1G (MT1G), mRNA  NM 006660  Homo sapiens metallothionein 1G (MT1G), mRNA  NM 006670  Homo sapiens superoxide dismutase 2, mitochondrial (SOD2), mRNA  NM 00668  Homo sapiens superoxide dismutase 2, mitochondrial (SOD2), mRNA  NM 000636  Homo sapiens superoxide dismutase 2, mitochondrial (SOD2), mRNA  NM 000637  Homo sapiens superoxide dismutase 2A (inducible, hepatocytes) (NOS2A), mRNA  NM 000629  Homo sapiens sitric oxide synthase 2A (inducible, hepatocytes) (NOS2A), mRNA  NM 000620  Homo sapiens interferon (alpha, beta and omega) receptor 1 (IFNAR1), mRNA  NM 000610  Homo sapiens solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2 (SLC11A2), mRNA  NM 000610  Homo sapiens CD4 antigen (PS5) (CD4)		
NM 021105 Homo sapiens phospholipid scramblase 1 (PLSCR1), mRNA NM 002957 Homo sapiens retinoid X receptor, alpha (RXRA), mRNA NM 006268 Homo sapiens requiem, apoptosis response zine finger gene (REQ), mRNA NM 001106 Homo sapiens activin A receptor, type IIB (ACVR2B), mRNA NM 001105 Homo sapiens activin A receptor, type IIB (ACVR2B), mRNA NM 001105 Homo sapiens activin A receptor, type II (ACVR1), mRNA NM 001570 Homo sapiens lectin, mannose-binding, 1 (LMAN1), mRNA NM 021083 Homo sapiens kell blood group precursor (McLeod phenotype) (XK), mRNA NM 012083 Homo sapiens kell blood group precursor (McLeod phenotype) (XK), mRNA NM 012083 Homo sapiens small proline-rich protein 2C (SPRR2C), mRNA NM 005570 Homo sapiens small proline-rich protein 2C (SPRR2C), mRNA NM 006518 Homo sapiens small proline-rich protein 2C (SPRR2C), mRNA NM 006550 Homo sapiens small proline-rich protein 2C (SPRR2C), mRNA NM 006553 Homo sapiens protein kinase, cGMP-dependent, type 1 (PRKG1), mRNA NM 00633 Homo sapiens protein kinase, cGMP-dependent, type 1 (PRKG1), mRNA NM 005987 Homo sapiens small proline-rich protein 1A (SPRR1A), mRNA NM 005987 Homo sapiens small proline-rich protein 1A (SPRR1A), mRNA NM 005990 Homo sapiens metallothionein 1X (MT1X), mRNA NM 005991 Homo sapiens metallothionein 1X (MT1X), mRNA NM 004618 Homo sapiens interleukin 18 binding protein (IL18BP), mRNA NM 004618 Homo sapiens superoxide dismutase 2, mitochondrial (SOD2), mRNA NM 000636 Homo sapiens superoxide dismutase 2, mitochondrial (SOD2), mRNA NM 000637 Homo sapiens interferon (alpha, beta and omega) receptor 1 (IFNAR1), mRNA NM 000639 Homo sapiens interferon (alpha, beta and omega) receptor 1 (IFNAR1), mRNA NM 000630 Homo sapiens interferon (alpha, beta and omega) receptor 1 (IFNAR1), mRNA NM 000631 Homo sapiens interferon (alpha, beta and omega) receptor 1 (IFNAR1), mRNA NM 000630 Homo sapiens interferon (alpha, beta and omega) receptor 1 (IFNAR1), mRNA NM 000610 Homo sapiens interferon (alpha, beta and omega) receptor 1 (IFNAR1), mRNA NM 000610 Homo sapiens SD-	14141_021090	
NM   006268   Homo sapiens retuinoid X receptor, alpha (RXRA), mRNA	NIM 021105	
NM 001106  Homo sapiens requiem, apoptosis response zinc finger gene (REQ), mRNA NM 001106  Homo sapiens activin A receptor, type IIB (ACVR2B), mRNA NM 001105  Homo sapiens activin A receptor, type III (ACVR2), mRNA NM 001105  Homo sapiens activin A receptor, type I (ACVR1), mRNA NM 001083  Homo sapiens lectin, mannose-binding, 1 (LMAN1), mRNA NM 021083  Homo sapiens kell blood group precursor (McLeod phenotype) (XK), mRNA NM 012083  Homo sapiens kell blood group precursor (McLeod phenotype) (XK), mRNA NM 006518  Homo sapiens small proline-rich protein 2C (SPRR2C), mRNA NM 006518  Homo sapiens small proline-rich protein 2C (SPRR2C), mRNA NM 006507  Homo sapiens small proline-rich protein 2C (SPRR2C), mRNA NM 006563  Homo sapiens Kruppel-like factor 1 (erythroid) (KLF1), mRNA NM 006258  Homo sapiens protein kinase, cGMP-dependent, type I (PRKG1), mRNA NM 006353  Homo sapiens protein kinase, cGMP-dependent, type I (PRKG1), mRNA NM 005987  Homo sapiens small proline-rich protein 1A (SPRR1A), mRNA NM 005987  Homo sapiens metallothionein 1X (MT1X), mRNA NM 005990  Homo sapiens metallothionein 1G (MT1G), mRNA NM 005699  Homo sapiens metallothionein 1G (MT1G), mRNA NM 004618  Homo sapiens advanced glycosylation end product-specific receptor (AGER), mRNA NM 00666  Homo sapiens glutathione reductase (GSR), mRNA NM 000636  Homo sapiens glutathione reductase (GSR), mRNA NM 000637  Homo sapiens interferon (alpha, beta and omega) receptor 1 (IFNAR1), mRNA NM 000636  Homo sapiens mitric oxide synthase 2A (inducible, hepatocytes) (NOS2A), mRNA NM 000629  Homo sapiens interferon (alpha, beta and omega) receptor 1 (IFNAR1), mRNA NM 000630  Homo sapiens interferon (alpha, beta and omega) receptor 1 (IFNAR1), mRNA NM 000630  Homo sapiens succear factor of kappa light polypeptide gene enhancer in B-cell 1 (p105) (NFKB1), mRNA NM 000637  Homo sapiens nitric oxide synthase 2A (inducible, hepatocytes) (NOS2A), mRNA NM 000630  Homo sapiens interferon, gamma (IFNG), mRNA NM 000617  Homo sapiens cD44 antigen (b55) (CD44), mRNA NM 000617		
NM 001616   Homo sapiens activin A receptor, type IIB (ACVR2B), mRNA   NM 001616   Homo sapiens activin A receptor, type II (ACVR2), mRNA   NM 0010570   Homo sapiens lectin, mannose-binding, 1 (LMAN1), mRNA   NM 021083   Homo sapiens Kell blood group precursor (McLeod phenotype) (XK), mRNA   NM 021083   Homo sapiens spoptosis-associated speck-like protein containing a CARD (ASC), mRNA   NM 013258   Homo sapiens small proline-rich protein 2C (SPRR2C), mRNA   NM 006518   Homo sapiens small proline-rich protein 2C (SPRR2C), mRNA   NM 006507   Homo sapiens regenerating islet-derived 1 beta (pancreatic stone protein, pancreatic thread protein) (REG1B), mRNA   NM 006508   Homo sapiens Kruppel-like factor 1 (erythroid) (KLF1), mRNA   NM 006538   Homo sapiens protein kinase, cGMP-dependent, type I (PRKG1), mRNA   NM 005987   Homo sapiens mall proline-rich protein 1A (SPRR1A), mRNA   NM 005950   Homo sapiens small proline-rich protein 1A (SPRR1A), mRNA   NM 005950   Homo sapiens metallothionein 1X (MT1X), mRNA   NM 005950   Homo sapiens metallothionein IG (MT1G), mRNA   NM 004618   Homo sapiens stopoisomerase (DNA) III alpha (TOP3A), mRNA   NM 0036609   Homo sapiens advanced glycosylation end product-specific receptor (AGER), mRNA   NM 000637   Homo sapiens glutathione reductase (GSR), mRNA   NM 000637   Homo sapiens strepulatory factor X, 2 (influences HLA class II expression) (RFX2), mRNA   NM 000639   Homo sapiens sitric oxide synthase 2 (inducible, hepatocytes) (NOS2A), mRNA   NM 000620   Homo sapiens interferon (alpha, beta and omega) receptor 1 (IFNAR1), mRNA   NM 000621   Homo sapiens nitric oxide synthase 2A (inducible, hepatocytes) (NOS2A), mRNA   NM 000621   Homo sapiens nitric oxide synthase 2A (inducible, hepatocytes) (NOS2A), mRNA   NM 000621   Homo sapiens nitric oxide synthase 2A (inducible, hepatocytes) (NOS2A), mRNA   NM 000617   Homo sapiens nitric oxide synthase 1 (neuronal) (NOS1), mRNA   NM 000617   Homo sapiens nitric oxide synthase 1 (neuronal) (NOS1), mRNA   NM 000610   Homo sapiens cD44 antige		
NM         001616         Homo sapiens activin A receptor, type I (ACVR2), mRNA           NM         001105         Homo sapiens activin A receptor, type I (ACVR1), mRNA           NM         005570         Homo sapiens Lectin, mannose-binding, I (LMANI), mRNA           NM         021083         Homo sapiens Rell blood group precursor (McLeod phenotype) (XK), mRNA           NM         013258         Homo sapiens apoptosis-associated speck-like protein containing a CARD (ASC), mRNA           NM         006518         Homo sapiens small proline-rich protein 2C (SPRR2C), mRNA           NM         006507         Homo sapiens small proline-rich protein 2C (EPRR2C), mRNA           NM         006563         Homo sapiens Kruppel-like factor 1 (erythroid) (KLF1), mRNA           NM         006563         Homo sapiens protein kinase, cGMP-dependent, type I (PRKGI), mRNA           NM         006528         Homo sapiens protein kinase, cGMP-dependent, type I (PRKGI), mRNA           NM         005987         Homo sapiens metallothionein 1X (MT1X), mRNA           NM         005987         Homo sapiens metallothionein 1X (MT1X), mRNA           NM         005950         Homo sapiens metallothionein 1G (MTIG), mRNA           NM         005950         Homo sapiens superoxide dismutase 2, micohondrial (SOD2), mRNA           NM         0006699         Homo sapiens superoxide dismutase 2,		
NM		
NM         005570         Homo sapiens lectin, mannose-binding, 1 (LMAN1), mRNA           NM         021083         Homo sapiens Kell blood group precursor (McLeod phenotype) (XK), mRNA           NM         013258         Homo sapiens apoptosis-associated speck-like protein containing a CARD (ASC), mRNA           NM         006518         Homo sapiens small proline-rich protein 2C (SPRR2C), mRNA           NM         006507         Homo sapiens small proline-rich protein 12C (SPRR2C), mRNA           NM         006563         Homo sapiens Kruppel-like factor 1 (erythroid) (KLF1), mRNA           NM         006563         Homo sapiens kruppel-like factor 1 (erythroid) (KLF1), mRNA           NM         006584         Homo sapiens protein kinase, cGMP-dependent, type 1 (PRKG1), mRNA           NM         006353         Homo sapiens protein kinase, cGMP-dependent, type 1 (PRKG1), mRNA           NM         005987         Homo sapiens small proline-rich protein 1A (SPR1A), mRNA           NM         005997         Homo sapiens metallothionein 1X (MT1X), mRNA           NM         005952         Homo sapiens metallothionein 1G (MT1G), mRNA           NM         005699         Homo sapiens interleukin 18 binding protein (IL18BP), mRNA           NM         001136         Homo sapiens stopoisomerase (DNA) III alpha (TOP3A), mRNA           NM         000636         Homo sapiens stopoi		
NM   013258		
NM_013258		
Mathematical Nation		
NM 006518   Homo sapiens small proline-rich protein 2C (SPRR2C), mRNA	14141_015256	
NM_006507	NM 006518	
pancreatic thread protein) (REG1B), mRNA  NM_006563 Homo sapiens Kruppel-like factor 1 (erythroid) (KLF1), mRNA  NM_006258 Homo sapiens protein kinase, cGMP-dependent, type I (PRKG1), mRNA  NM_006353 Homo sapiens high-mobility group (nonhistone chromosomal) protein 17-like 3 (HMG17L3), mRNA  NM_005987 Homo sapiens small proline-rich protein 1A (SPRR1A), mRNA  NM_005952 Homo sapiens metallothionein 1X (MT1X), mRNA  NM_005969 Homo sapiens metallothionein 1G (MT1G), mRNA  NM_001369 Homo sapiens interleukin 18 binding protein (IL18BP), mRNA  NM_00136 Homo sapiens advanced glycosylation end product-specific receptor (AGER), mRNA  NM_000866 Homo sapiens 5-hydroxytryptamine (serotonin) receptor 1F (HTR1F), mRNA  NM_000637 Homo sapiens glutathione reductase (GSR), mRNA  NM_000638 Homo sapiens superoxide dismutase 2, mitochondrial (SOD2), mRNA  NM_000639 Homo sapiens regulatory factor X, 2 (influences HLA class II expression) (RFX2), mRNA  NM_000629 Homo sapiens interferon (alpha, beta and omega) receptor 1 (IFNAR1), mRNA  NM_000629 Homo sapiens nuclear factor of kappa light polypeptide gene enhancer in B-cell 1 (p105) (NFKB1), mRNA  NM_000620 Homo sapiens interferon, gamma (IFNG), mRNA  NM_000610 Homo sapiens solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2 (SLC11A2), mRNA  NM_000616 Homo sapiens CD4 antigen (p55) (CD4), mRNA  NM_000610 Homo sapiens CD4 antigen (p55) (CD4), mRNA  NM_000610 Homo sapiens CD4 antigen (p55) (CD4), mRNA  NM_000610 Homo sapiens CD4 antigen (p55) (CD4), mRNA		
NM006563Homo sapiens Kruppel-like factor 1 (erythroid) (KLF1), mRNANM006258Homo sapiens protein kinase, cGMP-dependent, type I (PRKG1), mRNANM006353Homo sapiens high-mobility group (nonhistone chromosomal) protein 17-like 3 (HMG17L3), mRNANM005987Homo sapiens small proline-rich protein 1A (SPRR1A), mRNANM005950Homo sapiens metallothionein 1X (MT1X), mRNANM005950Homo sapiens metallothionein 1G (MT1G), mRNANM005699Homo sapiens interleukin 18 binding protein (IL18BP), mRNANM004618Homo sapiens topoisomerase (DNA) III alpha (TOP3A), mRNANM001136Homo sapiens 5-hydroxytryptamine (serotonin) receptor 1F (HTR1F), mRNANM000637Homo sapiens superoxide dismutase 2, mitochondrial (SOD2), mRNANM000636Homo sapiens superoxide dismutase 2, mitochondrial (SOD2), mRNANM000637Homo sapiens interferon (alpha, beta and omega) receptor 1 (IFNAR1), mRNANM000629Homo sapiens interferon (alpha, beta and omega) receptor 1 (IFNAR1), mRNANM000620Homo sapiens nuclear factor of kappa light polypeptide gene enhancer in B-cell 1 (p105) (NFKB1), mRNANM000621Homo sapiens solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2 (SLC11A2), mRNANM000617Homo sapiens cD4 antigen (p55) (CD4), mRNANM000610Homo sapiens CD5 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5, EJ16, EJ30, EL32 and G344) (CD59), mRNANM000610Homo sapiens CD44 antigen (homing function and Indian blood g	11111_000507	
NM_006258	NM 006563	Homo sapiens Kruppel-like factor 1 (erythroid) (KLF1), mRNA
NM_006353		
(HMG17L3), mRNA  NM 005987 Homo sapiens small proline-rich protein 1A (SPRR1A), mRNA  NM 005952 Homo sapiens metallothionein 1X (MT1X), mRNA  NM 005950 Homo sapiens metallothionein 1G (MT1G), mRNA  NM 005699 Homo sapiens interleukin 18 binding protein (IL18BP), mRNA  NM 004618 Homo sapiens topoisomerase (DNA) III alpha (TOP3A), mRNA  NM 001136 Homo sapiens advanced glycosylation end product-specific receptor (AGER), mRNA  NM 000866 Homo sapiens 5-hydroxytryptamine (serotonin) receptor 1F (HTR1F), mRNA  NM 000637 Homo sapiens glutathione reductase (GSR), mRNA  NM 000636 Homo sapiens superoxide dismutase 2, mitochondrial (SOD2), mRNA  NM 000635 Homo sapiens regulatory factor X, 2 (influences HLA class II expression)  (RFX2), mRNA  NM 000629 Homo sapiens interferon (alpha, beta and omega) receptor 1 (IFNAR1), mRNA  NM 000625 Homo sapiens nitric oxide synthase 2A (inducible, hepatocytes) (NOS2A),  mRNA  NM 000626 Homo sapiens factor of kappa light polypeptide gene enhancer in B-cell 1 (p105) (NFKB1), mRNA  NM 000621 Homo sapiens 5-hydroxytryptamine (serotonin) receptor 2A (HTR2A), mRNA  NM 000620 Homo sapiens interferon, gamma (IFNG), mRNA  NM 000610 Homo sapiens solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2 (SLC11A2), mRNA  NM 000616 Homo sapiens CD4 antigen (p55) (CD4), mRNA  NM 000610 Homo sapiens CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5, EJ16, EJ30, EL32 and G344) (CD59), mRNA  NM 000610 Homo sapiens CD44 antigen (homing function and Indian blood group system)		
NM_005987Homo sapiens small proline-rich protein 1A (SPRR1A), mRNANM_005952Homo sapiens metallothionein 1X (MT1X), mRNANM_005950Homo sapiens metallothionein 1G (MT1G), mRNANM_005699Homo sapiens interleukin 18 binding protein (IL18BP), mRNANM_004618Homo sapiens topoisomerase (DNA) III alpha (TOP3A), mRNANM_001136Homo sapiens advanced glycosylation end product-specific receptor (AGER), mRNANM_000866Homo sapiens 5-hydroxytryptamine (serotonin) receptor 1F (HTR1F), mRNANM_000637Homo sapiens glutathione reductase (GSR), mRNANM_000638Homo sapiens superoxide dismutase 2, mitochondrial (SOD2), mRNANM_000639Homo sapiens regulatory factor X, 2 (influences HLA class II expression) (RFX2), mRNANM_000629Homo sapiens interferon (alpha, beta and omega) receptor 1 (IFNAR1), mRNANM_000629Homo sapiens nuclear factor of kappa light polypeptide gene enhancer in B-cell 1 (p105) (NFKB1), mRNANM_003998Homo sapiens solute carrier factor of kappa light polypeptide gene enhancer in B-cell 1 (p105) (NFKB1), mRNANM_000610Homo sapiens solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2 (SLC11A2), mRNANM_000616Homo sapiens CD4 antigen (p55) (CD4), mRNANM_000610Homo sapiens CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5, EJ16, EJ30, EL32 and G344) (CD59), mRNANM_000610Homo sapiens CD44 antigen (homing function and Indian blood group system)	1111_000333	
NM005952Homo sapiens metallothionein 1X (MT1X), mRNANM005950Homo sapiens metallothionein 1G (MT1G), mRNANM005699Homo sapiens interleukin 18 binding protein (IL18BP), mRNANM004618Homo sapiens topoisomerase (DNA) III alpha (TOP3A), mRNANM001136Homo sapiens advanced glycosylation end product-specific receptor (AGER), mRNANM000866Homo sapiens 5-hydroxytryptamine (serotonin) receptor 1F (HTR1F), mRNANM000637Homo sapiens glutathione reductase (GSR), mRNANM000636Homo sapiens superoxide dismutase 2, mitochondrial (SOD2), mRNANM000635Homo sapiens regulatory factor X, 2 (influences HLA class II expression) (RFX2), mRNANM000629Homo sapiens interferon (alpha, beta and omega) receptor 1 (IFNAR1), mRNANM000629Homo sapiens nuclear factor of kappa light polypeptide gene enhancer in B-cell 1 (p105) (NFKB1), mRNANM003998Homo sapiens nuclear factor of kappa light polypeptide gene enhancer in B-cell 1 (p105) (NFKB1), mRNANM000620Homo sapiens interferon, gamma (IFNG), mRNANM000610Homo sapiens solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2 (SLC11A2), mRNANM000616Homo sapiens CD4 antigen (p55) (CD4), mRNANM000610Homo sapiens CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5, EJ16, EJ30, EL32 and G344) (CD59), mRNANM000610Homo sapiens CD44 antigen (homing function and Indian blood group system)	NM 005987	
NM005950Homo sapiens metallothionein 1G (MT1G), mRNANM005699Homo sapiens interleukin 18 binding protein (IL18BP), mRNANM004618Homo sapiens topoisomerase (DNA) III alpha (TOP3A), mRNANM001136Homo sapiens advanced glycosylation end product-specific receptor (AGER), mRNANM000866Homo sapiens 5-hydroxytryptamine (serotonin) receptor 1F (HTR1F), mRNANM000637Homo sapiens glutathione reductase (GSR), mRNANM000636Homo sapiens superoxide dismutase 2, mitochondrial (SOD2), mRNANM000635Homo sapiens regulatory factor X, 2 (influences HLA class II expression) (RFX2), mRNANM000629Homo sapiens interferon (alpha, beta and omega) receptor 1 (IFNAR1), mRNANM000629Homo sapiens nuclear factor of kappa light polypeptide gene enhancer in B-cell 1 (p105) (NFKB1), mRNANM003998Homo sapiens soluclear factor of kappa light polypeptide gene enhancer in B-cell 1 (p105) (NFKB1), mRNANM000621Homo sapiens f-hydroxytryptamine (serotonin) receptor 2A (HTR2A), mRNANM000620Homo sapiens interferon, gamma (IFNG), mRNANM000619Homo sapiens solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2 (SLC11A2), mRNANM000610Homo sapiens CD4 antigen (p55) (CD4), mRNANM000610Homo sapiens CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5, EJ16, EJ30, EL32 and G344) (CD59), mRNANM000610Homo sapiens CD44 antigen (homing function and Indian blood group system)		<u>, , , , , , , , , , , , , , , , , , , </u>
NM_005699Homo sapiens interleukin 18 binding protein (IL18BP), mRNANM_004618Homo sapiens topoisomerase (DNA) III alpha (TOP3A), mRNANM_001136Homo sapiens advanced glycosylation end product-specific receptor (AGER), mRNANM_000866Homo sapiens 5-hydroxytryptamine (serotonin) receptor 1F (HTR1F), mRNANM_000637Homo sapiens glutathione reductase (GSR), mRNANM_000638Homo sapiens superoxide dismutase 2, mitochondrial (SOD2), mRNANM_000639Homo sapiens regulatory factor X, 2 (influences HLA class II expression) (RFX2), mRNANM_000629Homo sapiens interferon (alpha, beta and omega) receptor 1 (IFNAR1), mRNANM_000625Homo sapiens nitric oxide synthase 2A (inducible, hepatocytes) (NOS2A), mRNANM_003998Homo sapiens nuclear factor of kappa light polypeptide gene enhancer in B-cell 1 (p105) (NFKB1), mRNANM_000621Homo sapiens 5-hydroxytryptamine (serotonin) receptor 2A (HTR2A), mRNANM_000620Homo sapiens nitric oxide synthase 1 (neuronal) (NOS1), mRNANM_000619Homo sapiens interferon, gamma (IFNG), mRNANM_000610Homo sapiens CD4 antigen (p55) (CD4), mRNANM_000611Homo sapiens CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5, EJ16, EJ30, EL32 and G344) (CD59), mRNANM_000610Homo sapiens CD44 antigen (homing function and Indian blood group system)		
NM004618Homo sapiens topoisomerase (DNA) III alpha (TOP3A), mRNANM_001136Homo sapiens advanced glycosylation end product-specific receptor (AGER), mRNANM000866Homo sapiens 5-hydroxytryptamine (serotonin) receptor 1F (HTR1F), mRNANM000637Homo sapiens glutathione reductase (GSR), mRNANM000636Homo sapiens superoxide dismutase 2, mitochondrial (SOD2), mRNANM_000635Homo sapiens regulatory factor X, 2 (influences HLA class II expression) (RFX2), mRNANM000629Homo sapiens interferon (alpha, beta and omega) receptor 1 (IFNAR1), mRNANM_000625Homo sapiens nitric oxide synthase 2A (inducible, hepatocytes) (NOS2A), mRNANM_003998Homo sapiens nuclear factor of kappa light polypeptide gene enhancer in B-cell 1 (p105) (NFKB1), mRNANM000621Homo sapiens 5-hydroxytryptamine (serotonin) receptor 2A (HTR2A), mRNANM000620Homo sapiens interferon, gamma (IFNG), mRNANM000619Homo sapiens solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2 (SLC11A2), mRNANM000616Homo sapiens CD4 antigen (p55) (CD4), mRNANM_000611Homo sapiens CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5, EJ16, EJ30, EL32 and G344) (CD59), mRNANM_000610Homo sapiens CD44 antigen (homing function and Indian blood group system)		
NM_001136 Homo sapiens advanced glycosylation end product-specific receptor (AGER), mRNA  NM_000866 Homo sapiens 5-hydroxytryptamine (serotonin) receptor 1F (HTR1F), mRNA  NM_000637 Homo sapiens glutathione reductase (GSR), mRNA  NM_000636 Homo sapiens superoxide dismutase 2, mitochondrial (SOD2), mRNA  NM_000635 Homo sapiens regulatory factor X, 2 (influences HLA class II expression) (RFX2), mRNA  NM_000629 Homo sapiens interferon (alpha, beta and omega) receptor 1 (IFNAR1), mRNA  NM_000625 Homo sapiens nitric oxide synthase 2A (inducible, hepatocytes) (NOS2A), mRNA  NM_003998 Homo sapiens nuclear factor of kappa light polypeptide gene enhancer in B-cell 1 (p105) (NFKB1), mRNA  NM_000621 Homo sapiens 5-hydroxytryptamine (serotonin) receptor 2A (HTR2A), mRNA  NM_000620 Homo sapiens interferon, gamma (IFNG), mRNA  NM_000619 Homo sapiens solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2 (SLC11A2), mRNA  NM_000616 Homo sapiens CD4 antigen (p55) (CD4), mRNA  NM_000611 Homo sapiens CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5, EJ16, EJ30, EL32 and G344) (CD59), mRNA  NM_000610 Homo sapiens CD44 antigen (homing function and Indian blood group system)		
mRNA  NM_000866 Homo sapiens 5-hydroxytryptamine (serotonin) receptor 1F (HTR1F), mRNA  NM_000637 Homo sapiens glutathione reductase (GSR), mRNA  NM_000636 Homo sapiens superoxide dismutase 2, mitochondrial (SOD2), mRNA  NM_000635 Homo sapiens regulatory factor X, 2 (influences HLA class II expression)  (RFX2), mRNA  NM_000629 Homo sapiens interferon (alpha, beta and omega) receptor 1 (IFNAR1), mRNA  NM_000625 Homo sapiens nitric oxide synthase 2A (inducible, hepatocytes) (NOS2A),  mRNA  NM_003998 Homo sapiens nuclear factor of kappa light polypeptide gene enhancer in B-cell  1 (p105) (NFKB1), mRNA  NM_000621 Homo sapiens 5-hydroxytryptamine (serotonin) receptor 2A (HTR2A), mRNA  NM_000620 Homo sapiens nitric oxide synthase 1 (neuronal) (NOS1), mRNA  NM_000619 Homo sapiens solute carrier family 11 (proton-coupled divalent metal ion  transporters), member 2 (SLC11A2), mRNA  NM_000616 Homo sapiens CD4 antigen (p55) (CD4), mRNA  NM_000610 Homo sapiens CD44 antigen (homing function and Indian blood group system)  NM_000610 Homo sapiens CD44 antigen (homing function and Indian blood group system)		
NM_000636 Homo sapiens 5-hydroxytryptamine (serotonin) receptor 1F (HTR1F), mRNA NM_000637 Homo sapiens glutathione reductase (GSR), mRNA NM_000636 Homo sapiens superoxide dismutase 2, mitochondrial (SOD2), mRNA NM_000635 Homo sapiens regulatory factor X, 2 (influences HLA class II expression) (RFX2), mRNA NM_000629 Homo sapiens interferon (alpha, beta and omega) receptor 1 (IFNAR1), mRNA NM_000625 Homo sapiens nitric oxide synthase 2A (inducible, hepatocytes) (NOS2A), mRNA NM_003998 Homo sapiens nuclear factor of kappa light polypeptide gene enhancer in B-cell 1 (p105) (NFKB1), mRNA NM_000621 Homo sapiens 5-hydroxytryptamine (serotonin) receptor 2A (HTR2A), mRNA NM_000620 Homo sapiens nitric oxide synthase 1 (neuronal) (NOS1), mRNA NM_000619 Homo sapiens interferon, gamma (IFNG), mRNA NM_000617 Homo sapiens solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2 (SLC11A2), mRNA NM_000616 Homo sapiens CD4 antigen (p55) (CD4), mRNA NM_000611 Homo sapiens CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5, EJ16, EJ30, EL32 and G344) (CD59), mRNA NM_000610 Homo sapiens CD44 antigen (homing function and Indian blood group system)		
NM_000637 Homo sapiens glutathione reductase (GSR), mRNA  NM_000636 Homo sapiens superoxide dismutase 2, mitochondrial (SOD2), mRNA  NM_000635 Homo sapiens regulatory factor X, 2 (influences HLA class II expression) (RFX2), mRNA  NM_000629 Homo sapiens interferon (alpha, beta and omega) receptor 1 (IFNAR1), mRNA  NM_000625 Homo sapiens nitric oxide synthase 2A (inducible, hepatocytes) (NOS2A), mRNA  NM_003998 Homo sapiens nuclear factor of kappa light polypeptide gene enhancer in B-cell 1 (p105) (NFKB1), mRNA  NM_000621 Homo sapiens 5-hydroxytryptamine (serotonin) receptor 2A (HTR2A), mRNA  NM_000620 Homo sapiens nitric oxide synthase 1 (neuronal) (NOS1), mRNA  NM_000619 Homo sapiens interferon, gamma (IFNG), mRNA  NM_000617 Homo sapiens solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2 (SLC11A2), mRNA  NM_000616 Homo sapiens CD4 antigen (p55) (CD4), mRNA  NM_000610 Homo sapiens CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5, EJ16, EJ30, EL32 and G344) (CD59), mRNA  NM_000610 Homo sapiens CD44 antigen (homing function and Indian blood group system)	NM 000866	Homo sapiens 5-hydroxytryptamine (serotonin) receptor 1F (HTR1F), mRNA
NM_000635 Homo sapiens regulatory factor X, 2 (influences HLA class II expression) (RFX2), mRNA  NM_000629 Homo sapiens interferon (alpha, beta and omega) receptor 1 (IFNAR1), mRNA  NM_000625 Homo sapiens nitric oxide synthase 2A (inducible, hepatocytes) (NOS2A), mRNA  NM_003998 Homo sapiens nuclear factor of kappa light polypeptide gene enhancer in B-cell 1 (p105) (NFKB1), mRNA  NM_000621 Homo sapiens 5-hydroxytryptamine (serotonin) receptor 2A (HTR2A), mRNA  NM_000620 Homo sapiens nitric oxide synthase 1 (neuronal) (NOS1), mRNA  NM_000619 Homo sapiens interferon, gamma (IFNG), mRNA  NM_000617 Homo sapiens solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2 (SLC11A2), mRNA  NM_000616 Homo sapiens CD4 antigen (p55) (CD4), mRNA  NM_000610 Homo sapiens CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5, EJ16, EJ30, EL32 and G344) (CD59), mRNA  NM_000610 Homo sapiens CD44 antigen (homing function and Indian blood group system)	NM 000637	Homo sapiens glutathione reductase (GSR), mRNA
(RFX2), mRNA  NM_000629 Homo sapiens interferon (alpha, beta and omega) receptor 1 (IFNAR1), mRNA  NM_000625 Homo sapiens nitric oxide synthase 2A (inducible, hepatocytes) (NOS2A), mRNA  NM_003998 Homo sapiens nuclear factor of kappa light polypeptide gene enhancer in B-cell 1 (p105) (NFKB1), mRNA  NM_000621 Homo sapiens 5-hydroxytryptamine (serotonin) receptor 2A (HTR2A), mRNA  NM_000620 Homo sapiens nitric oxide synthase 1 (neuronal) (NOS1), mRNA  NM_000619 Homo sapiens interferon, gamma (IFNG), mRNA  NM_000617 Homo sapiens solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2 (SLC11A2), mRNA  NM_000616 Homo sapiens CD4 antigen (p55) (CD4), mRNA  NM_000611 Homo sapiens CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5, EJ16, EJ30, EL32 and G344) (CD59), mRNA  NM_000610 Homo sapiens CD44 antigen (homing function and Indian blood group system)	NM 000636	Homo sapiens superoxide dismutase 2, mitochondrial (SOD2), mRNA
(RFX2), mRNA  NM_000629 Homo sapiens interferon (alpha, beta and omega) receptor 1 (IFNAR1), mRNA  NM_000625 Homo sapiens nitric oxide synthase 2A (inducible, hepatocytes) (NOS2A), mRNA  NM_003998 Homo sapiens nuclear factor of kappa light polypeptide gene enhancer in B-cell 1 (p105) (NFKB1), mRNA  NM_000621 Homo sapiens 5-hydroxytryptamine (serotonin) receptor 2A (HTR2A), mRNA  NM_000620 Homo sapiens nitric oxide synthase 1 (neuronal) (NOS1), mRNA  NM_000619 Homo sapiens interferon, gamma (IFNG), mRNA  NM_000617 Homo sapiens solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2 (SLC11A2), mRNA  NM_000616 Homo sapiens CD4 antigen (p55) (CD4), mRNA  NM_000611 Homo sapiens CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5, EJ16, EJ30, EL32 and G344) (CD59), mRNA  NM_000610 Homo sapiens CD44 antigen (homing function and Indian blood group system)	NM 000635	Homo sapiens regulatory factor X, 2 (influences HLA class II expression)
NM_000625 Homo sapiens nitric oxide synthase 2A (inducible, hepatocytes) (NOS2A), mRNA  NM_003998 Homo sapiens nuclear factor of kappa light polypeptide gene enhancer in B-cell 1 (p105) (NFKB1), mRNA  NM_000621 Homo sapiens 5-hydroxytryptamine (serotonin) receptor 2A (HTR2A), mRNA  NM_000620 Homo sapiens nitric oxide synthase 1 (neuronal) (NOS1), mRNA  NM_000619 Homo sapiens interferon, gamma (IFNG), mRNA  NM_000617 Homo sapiens solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2 (SLC11A2), mRNA  NM_000616 Homo sapiens CD4 antigen (p55) (CD4), mRNA  NM_000610 Homo sapiens CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5, EJ16, EJ30, EL32 and G344) (CD59), mRNA  NM_000610 Homo sapiens CD44 antigen (homing function and Indian blood group system)		
mRNA  NM_003998 Homo sapiens nuclear factor of kappa light polypeptide gene enhancer in B-cell 1 (p105) (NFKB1), mRNA  NM_000621 Homo sapiens 5-hydroxytryptamine (serotonin) receptor 2A (HTR2A), mRNA  NM_000620 Homo sapiens nitric oxide synthase 1 (neuronal) (NOS1), mRNA  NM_000619 Homo sapiens interferon, gamma (IFNG), mRNA  NM_000617 Homo sapiens solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2 (SLC11A2), mRNA  NM_000616 Homo sapiens CD4 antigen (p55) (CD4), mRNA  NM_000611 Homo sapiens CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5, EJ16, EJ30, EL32 and G344) (CD59), mRNA  NM_000610 Homo sapiens CD44 antigen (homing function and Indian blood group system)	NM_000629	
NM_003998 Homo sapiens nuclear factor of kappa light polypeptide gene enhancer in B-cell 1 (p105) (NFKB1), mRNA  NM_000621 Homo sapiens 5-hydroxytryptamine (serotonin) receptor 2A (HTR2A), mRNA  NM_000620 Homo sapiens nitric oxide synthase 1 (neuronal) (NOS1), mRNA  NM_000619 Homo sapiens interferon, gamma (IFNG), mRNA  NM_000617 Homo sapiens solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2 (SLC11A2), mRNA  NM_000616 Homo sapiens CD4 antigen (p55) (CD4), mRNA  NM_000611 Homo sapiens CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5, EJ16, EJ30, EL32 and G344) (CD59), mRNA  NM_000610 Homo sapiens CD44 antigen (homing function and Indian blood group system)	NM_000625	
1 (p105) (NFKB1), mRNA  NM_000621 Homo sapiens 5-hydroxytryptamine (serotonin) receptor 2A (HTR2A), mRNA  NM_000620 Homo sapiens nitric oxide synthase 1 (neuronal) (NOS1), mRNA  NM_000619 Homo sapiens interferon, gamma (IFNG), mRNA  NM_000617 Homo sapiens solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2 (SLC11A2), mRNA  NM_000616 Homo sapiens CD4 antigen (p55) (CD4), mRNA  NM_000611 Homo sapiens CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5, EJ16, EJ30, EL32 and G344) (CD59), mRNA  NM_000610 Homo sapiens CD44 antigen (homing function and Indian blood group system)	NM 003998	Homo sapiens nuclear factor of kappa light polypeptide gene enhancer in B-cells
NM_000621Homo sapiens 5-hydroxytryptamine (serotonin) receptor 2A (HTR2A), mRNANM_000620Homo sapiens nitric oxide synthase 1 (neuronal) (NOS1), mRNANM_000619Homo sapiens interferon, gamma (IFNG), mRNANM_000617Homo sapiens solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2 (SLC11A2), mRNANM_000616Homo sapiens CD4 antigen (p55) (CD4), mRNANM_000611Homo sapiens CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5, EJ16, EJ30, EL32 and G344) (CD59), mRNANM_000610Homo sapiens CD44 antigen (homing function and Indian blood group system)	_	
NM_000620Homo sapiens nitric oxide synthase 1 (neuronal) (NOS1), mRNANM_000619Homo sapiens interferon, gamma (IFNG), mRNANM_000617Homo sapiens solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2 (SLC11A2), mRNANM_000616Homo sapiens CD4 antigen (p55) (CD4), mRNANM_000611Homo sapiens CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5, EJ16, EJ30, EL32 and G344) (CD59), mRNANM_000610Homo sapiens CD44 antigen (homing function and Indian blood group system)	NM_000621	
NM_000619Homo sapiens interferon, gamma (IFNG), mRNANM_000617Homo sapiens solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2 (SLC11A2), mRNANM_000616Homo sapiens CD4 antigen (p55) (CD4), mRNANM_000611Homo sapiens CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5, EJ16, EJ30, EL32 and G344) (CD59), mRNANM_000610Homo sapiens CD44 antigen (homing function and Indian blood group system)		
NM_000617Homo sapiens solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2 (SLC11A2), mRNANM_000616Homo sapiens CD4 antigen (p55) (CD4), mRNANM_000611Homo sapiens CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5, EJ16, EJ30, EL32 and G344) (CD59), mRNANM_000610Homo sapiens CD44 antigen (homing function and Indian blood group system)	NM_000619	
transporters), member 2 (SLC11A2), mRNA  NM_000616 Homo sapiens CD4 antigen (p55) (CD4), mRNA  NM_000611 Homo sapiens CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5, EJ16, EJ30, EL32 and G344) (CD59), mRNA  NM_000610 Homo sapiens CD44 antigen (homing function and Indian blood group system)		
NM_000611 Homo sapiens CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5, EJ16, EJ30, EL32 and G344) (CD59), mRNA  NM_000610 Homo sapiens CD44 antigen (homing function and Indian blood group system)		transporters), member 2 (SLC11A2), mRNA
antibodies 16.3A5, EJ16, EJ30, EL32 and G344) (CD59), mRNA  NM_000610 Homo sapiens CD44 antigen (homing function and Indian blood group system)	NM_000616	
NM_000610 Homo sapiens CD44 antigen (homing function and Indian blood group system)	NM_000611	Homo sapiens CD59 antigen p18-20 (antigen identified by monoclonal
•	NM_000610	
(CD44), mRNA		
NM_000603 Homo sapiens nitric oxide synthase 3 (endothelial cell) (NOS3), mRNA		
NM_000597 Homo sapiens insulin-like growth factor binding protein 2 (36kD) (IGFBP2),	NM_000597	
mRNA		
NM_000594 Homo sapiens tumor necrosis factor (TNF superfamily, member 2) (TNF),	NM_000594	
mRNA		mRNA

omo sapiens interleukin 15 (IL15), mRNA omo sapiens interleukin 2 (IL2), mRNA omo sapiens interleukin 1 receptor antagonist (IL1RN), mRNA omo sapiens interleukin 1, beta (IL1B), mRNA omo sapiens decay accelerating factor for complement (CD55, Cromer blood oup system) (DAF), mRNA omo sapiens interleukin 10 (IL10), mRNA omo sapiens Fc fragment of IgG, low affinity IIIb, receptor for (CD16) CCGR3B), mRNA omo sapiens C-reactive protein, pentraxin-related (CRP), mRNA omo sapiens Fc fragment of IgG, high affinity Ia, receptor for (CD64) GCGR1A), mRNA
omo sapiens interleukin 1 receptor antagonist (IL1RN), mRNA omo sapiens interleukin 1, beta (IL1B), mRNA omo sapiens decay accelerating factor for complement (CD55, Cromer blood oup system) (DAF), mRNA omo sapiens interleukin 10 (IL10), mRNA omo sapiens Fc fragment of IgG, low affinity IIIb, receptor for (CD16) FCGR3B), mRNA omo sapiens C-reactive protein, pentraxin-related (CRP), mRNA tomo sapiens Fc fragment of IgG, high affinity Ia, receptor for (CD64) FCGR1A), mRNA
omo sapiens interleukin 1, beta (IL1B), mRNA omo sapiens decay accelerating factor for complement (CD55, Cromer blood oup system) (DAF), mRNA omo sapiens interleukin 10 (IL10), mRNA omo sapiens Fc fragment of IgG, low affinity IIIb, receptor for (CD16) FCGR3B), mRNA omo sapiens C-reactive protein, pentraxin-related (CRP), mRNA omo sapiens Fc fragment of IgG, high affinity Ia, receptor for (CD64) FCGR1A), mRNA
omo sapiens decay accelerating factor for complement (CD55, Cromer blood coup system) (DAF), mRNA omo sapiens interleukin 10 (IL10), mRNA omo sapiens Fc fragment of IgG, low affinity IIIb, receptor for (CD16) FCGR3B), mRNA omo sapiens C-reactive protein, pentraxin-related (CRP), mRNA omo sapiens Fc fragment of IgG, high affinity Ia, receptor for (CD64) FCGR1A), mRNA
oup system) (DAF), mRNA omo sapiens interleukin 10 (IL10), mRNA omo sapiens Fc fragment of IgG, low affinity IIIb, receptor for (CD16) FCGR3B), mRNA omo sapiens C-reactive protein, pentraxin-related (CRP), mRNA omo sapiens Fc fragment of IgG, high affinity Ia, receptor for (CD64) FCGR1A), mRNA
omo sapiens interleukin 10 (IL10), mRNA omo sapiens Fc fragment of IgG, low affinity IIIb, receptor for (CD16) FCGR3B), mRNA omo sapiens C-reactive protein, pentraxin-related (CRP), mRNA omo sapiens Fc fragment of IgG, high affinity Ia, receptor for (CD64) FCGR1A), mRNA
omo sapiens Fc fragment of IgG, low affinity IIIb, receptor for (CD16) FCGR3B), mRNA omo sapiens C-reactive protein, pentraxin-related (CRP), mRNA omo sapiens Fc fragment of IgG, high affinity Ia, receptor for (CD64) FCGR1A), mRNA
CCGR3B), mRNA  Tomo sapiens C-reactive protein, pentraxin-related (CRP), mRNA  Tomo sapiens Fc fragment of IgG, high affinity Ia, receptor for (CD64)  FCGR1A), mRNA
omo sapiens C-reactive protein, pentraxin-related (CRP), mRNA omo sapiens Fc fragment of IgG, high affinity Ia, receptor for (CD64) FCGR1A), mRNA
omo sapiens Fc fragment of IgG, high affinity Ia, receptor for (CD64) FCGR1A), mRNA
FCGR1A), mRNA
omo sapiens interleukin 5 receptor, alpha (IL5RA), mRNA
Iomo saniens glutathione S-transferase M1 (GSTM1), mRNA
Iomo sapiens doublecortex; lissencephaly, X-linked (doublecortin) (DCX),
Iomo sapiens pyruvate kinase, liver and RBC (PKLR), nuclear gene encoding
uitochondrial protein, mRNA
Jomo sanjens myosin VA (heavy polypeptide 12, myoxin) (MYO5A), mRNA
Iomo sapiens potassium inwardly-rectifying channel, subfamily J, member 11
KCNJ11), mRNA
Homo sapiens myotubularin related protein 3 (MTMR3), mRNA
Homo sapiens neuromedin B (NMB), mRNA
Iomo sapiens interferon, alpha 4 (IFNA4), mRNA
Homo sapiens serum amyloid A4, constitutive (SAA4), mRNA
Homo sapiens pituitary tumor-transforming 2 (PTTG2), mRNA
Homo sapiens NADH dehydrogenase (ubiquinone) flavoprotein 3 (10kD)
NDUFV3), mRNA
Homo sapiens metallothionein 1H (MT1H), mRNA
Homo sapiens retinoschisis (X-linked, juvenile) 1 (RS1), mRNA
Homo sapiens nuclear factor I/C (CCAAT-binding transcription factor) (NFIC),
mRNA
Homo sapiens gap junction protein, beta 5 (connexin 31.1) (GJB5), mRNA
Homo sapiens cofactor required for Sp1 transcriptional activation, subunit 6 (77kD) (CRSP6), mRNA
Homo sapiens CD74 antigen (invariant polypeptide of major histocompatibility
complex, class II antigen-associated) (CD74), mRNA
Homo sapiens protein kinase, Y-linked (PRKY), mRNA
Homo sapiens nucleophosmin (nucleolar phosphoprotein B23, numatrin) (NPM1), mRNA
Homo sapiens inhibitor of DNA binding 3, dominant negative helix-loop-helix protein (ID3), mRNA
Homo sapiens farnesyltransferase, CAAX box, beta (FNTB), mRNA
Homo sapiens N-acetyltransferase, homolog of S. cerevisiae ARD1 (ARD1),
mRNA
Homo sapiens CD19 antigen (CD19), mRNA
Homo sapiens ras homolog gene family, member A (ARHA), mRNA
Homo saniens sarcoglycan, epsilon (SGCE), mRNA
Homo sapiens tumor necrosis factor receptor superfamily, member 10c, decoy
without an intracellular domain (TNFRSF10C), mRNA
Homo sapiens zinc finger protein 202 (ZNF202), mRNA
Homo sapiens zinc finger protein 189 (ZNF189), mRNA

NM 003316	Homo ganiana totratricamentida repost domain 2 (TTC2) TNIA
NM 003166	Homo sapiens tetratricopeptide repeat domain 3 (TTC3), mRNA
1003100	Homo sapiens sulfotransferase family, cytosolic, 1A, phenol-preferring, member 3 (SULT1A3), mRNA
NM 003117	Homo sapiens sperm adhesion molecule 1 (PH-20 hyaluronidase, zona pellucida
11111_003117	binding) (SPAM1), mRNA
NM 002222	Homo sapiens inositol 1,4,5-triphosphate receptor, type 1 (ITPR1), mRNA
NM 001532	Homo sapiens solute carrier family 29 (nucleoside transporters), member 2
1111_001052	(SLC29A2), mRNA
NM 001437	Homo sapiens estrogen receptor 2 (ER beta) (ESR2), mRNA
NM 001331	Homo sapiens catenin (cadherin-associated protein), delta 1 (CTNND1), mRNA
NM 001307	Homo sapiens claudin 7 (CLDN7), mRNA
NM 001194	Homo sapiens hyperpolarization activated cyclic nucleotide-gated potassium
_	channel 2 (HCN2), mRNA
NM 001175	Homo sapiens Rho GDP dissociation inhibitor (GDI) beta (ARHGDIB), mRNA
NM_000936	Homo sapiens pancreatic lipase (PNLIP), mRNA
NM_000641	Homo sapiens interleukin 11 (IL11), mRNA
NM_000640	Homo sapiens interleukin 13 receptor, alpha 2 (IL13RA2), mRNA
NM_000615	Homo sapiens neural cell adhesion molecule 1 (NCAM1), mRNA
NM_000609	Homo sapiens stromal cell-derived factor 1 (SDF1), mRNA
NM_000600	Homo sapiens interleukin 6 (interferon, beta 2) (IL6), mRNA
NM_000599	Homo sapiens insulin-like growth factor binding protein 5 (IGFBP5), mRNA
NM_000590	Homo sapiens interleukin 9 (IL9), mRNA
NM_000584	Homo sapiens interleukin 8 (IL8), mRNA
NM_000581	Homo sapiens glutathione peroxidase 1 (GPX1), mRNA
NM_000560	Homo sapiens CD53 antigen (CD53), mRNA
NM_000528	Homo sapiens mannosidase, alpha, class 2B, member 1 (MAN2B1), mRNA
NM_000404	Homo sapiens galactosidase, beta 1 (GLB1), mRNA
NM_001275	Homo sapiens chromogranin A (parathyroid secretory protein 1) (CHGA), mRNA
NM_006768	Homo sapiens BRCA1 associated protein (BRAP), mRNA
NM_003469	Homo sapiens secretogranin II (chromogranin C) (SCG2), mRNA
NM_012326	Homo sapiens microtubule-associated protein, RP/EB family, member 3
	(MAPRE3), mRNA
NM_021057	Homo sapiens interferon, alpha 7 (IFNA7), mRNA
NM_021062	Homo sapiens H2B histone family, member F (H2BFF), mRNA
NM_021063	Homo sapiens H2B histone family, member B (H2BFB), mRNA
NM_021065	Homo sapiens H2A histone family, member G (H2AFG), mRNA
NM_004146	Homo sapiens NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 7 (18kD,
ND4 00:545	B18) (NDUFB7), mRNA
NM_001746	Homo sapiens calnexin (CANX), mRNA
NM_003661	Homo sapiens apolipoprotein L (APOL), mRNA
NM_021052	Homo sapiens H2A histone family, member A (H2AFA), mRNA
NM_020988	Homo sapiens guanine nucleotide binding protein (G protein), alpha activating activity polypeptide O (GNAO1), mRNA
NM 000133	Homo sapiens coagulation factor IX (plasma thromboplastic component,
	Christmas disease, hemophilia B) (F9), mRNA
NM_000130	Homo sapiens coagulation factor V (proaccelerin, labile factor) (F5), mRNA
NM_001993	Homo sapiens coagulation factor III (thromboplastin, tissue factor) (F3), mRNA
NM_020689	Homo sapiens sodium calcium exchanger (NCKX3), mRNA
NM_021033	Homo sapiens RAP2A, member of RAS oncogene family (RAP2A), mRNA
NM_021023	Homo sapiens complement factor H related 3 (FHR-3), mRNA
NM_021026	Homo sapiens ret finger protein-like 1 (RFPL1), mRNA

NM_021008	Homo sapiens suppressin (nuclear deformed epidermal autoregulatory factor-1
NB ( 020002	(DEAF-1)-related) (SPN), mRNA  Homo sapiens B-cell CLL/lymphoma 7A (BCL7A), mRNA
NM_020993	Homo sapiens B-cen CLL/lymphoma /A (BCL/A), mRNA  Homo sapiens cancer/testis antigen 2 (CTAG2), mRNA
NM_020994	Homo sapiens pituitary tumor-transforming 3 (PTTG3), mRNA
NM_021000	
NM_020997	Homo sapiens left-right determination, factor B (LEFTB), mRNA
NM_021014	Homo sapiens synovial sarcoma, X breakpoint 3 (SSX3), mRNA
NM_021015	Homo sapiens synovial sarcoma, X breakpoint 5 (SSX5), mRNA
NM_021007	Homo sapiens sodium channel, voltage-gated, type II, alpha 2 polypeptide (SCN2A2), mRNA
NM_021012	Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 12 (KCNJ12), mRNA
NM 020995	Homo sapiens haptoglobin-related protein (HPR), mRNA
NM_000347	Homo sapiens spectrin, beta, erythrocytic (includes spherocytosis, clinical type I) (SPTB), mRNA
NM 007032	Homo sapiens putative nuclear protein (HRIHFB2122), mRNA
NM 001320	Homo sapiens casein kinase 2, beta polypeptide (CSNK2B), mRNA
NM 013252	Homo sapiens C-type (calcium dependent, carbohydrate-recognition domain)
11112_013232	lectin, superfamily member 5 (CLECSF5), mRNA
NM 020978	Homo sapiens amylase, alpha 2B; pancreatic (AMY2B), mRNA
NM 020636	Homo sapiens zinc finger protein 275 (ZNF275), mRNA
NM 020547	Homo sapiens anti-Mullerian hormone receptor, type II (AMHR2), mRNA
NM 020974	Homo sapiens CEGP1 protein (CEGP1), mRNA
NM 020681	Homo sapiens HT018 protein (HT018), mRNA
NM 020676	Homo sapiens lipase protein (LOC57406), mRNA
NM 020672	Homo sapiens S100-type calcium binding protein A14 (LOC57402), mRNA
NM 020661	Homo sapiens activation-induced cytidine deaminase (AICDA), mRNA
NM 020657	Homo sapiens zinc finger protein 304 (ZNF304), mRNA
NM 020654	Homo sapiens sentrin/SUMO-specific protease (SENP7), mRNA
NM 020646	Homo sapiens reserved (ASCL3), mRNA
NM 020640	Homo sapiens RP42 homolog (RP42), mRNA
	Homo sapiens ankyrin repeat domain 3 (ANKRD3), mRNA
NM_020639	Homo sapiens ATPase, H(+)-transporting, lysosomal, noncatalytic accessory
NM_020632	
ND 4 020649	protein 1B (ATP6N1B), mRNA
NM_020648	Homo sapiens twisted gastrulation (TSG), mRNA
NM_018970	Homo sapiens G protein-coupled receptor 85 (GPR85), mRNA
NM_003901	Homo sapiens sphingosine-1-phosphate lyase 1 (SGPL1), mRNA
NM_014292	Homo sapiens chromobox homolog 6 (CBX6), mRNA
NM_006735	Homo sapiens homeo box A2 (HOXA2), mRNA
NM_019041	Homo sapiens similar to prokaryotic-type class I peptide chain release factors (LOC54516), mRNA
NM_014428	Homo sapiens tight junction protein 3 (zona occludens 3) (TJP3), mRNA
NM_020466	Homo sapiens hypothetical protein dJ122O8.2 (DJ122O8.2), mRNA
NM_020448	Homo sapiens hypothetical protein dJ462O23.2 (DJ462O23.2), mRNA
NM_020425	Homo sapiens hypothetical protein DKFZp586E1923 (DKFZP586E1923), mRNA
NM_020424	Homo sapiens hypothetical protein A-211C6.1 (LOC57149), mRNA
NM 020317	Homo sapiens hypothetical protein dJ465N24.2.1 (DJ465N24.2.1), mRNA
NM 020315	Homo sapiens hypothetical protein dJ37E16.5 (DJ37E16.5), mRNA
NM 020313	Homo sapiens hypothetical protein (LOC57019), mRNA
NM_019897	Homo sapiens olfactory receptor, family 2, subfamily S, member 2 (OR2S2),
	mRNA
L	14444

NM 019605	Homo sapiens hypothetical protein (DJ667H12.2), mRNA
NM 019601	Homo sapiens Sushi domain (SCR repeat) containing (BK65A6.2), mRNA
NM 018433	Homo sapiens putative zinc finger protein (LOC55818), mRNA
NM 019095	Homo sapiens hypothetical protein (LOC54675), mRNA
NM 019089	Homo sapiens hairy and enhancer of split (Drosophila) homolog 2 (HES2),
11111	mRNA
NM 018982	Homo sapiens hypothetical protein (DJ167A19.1), mRNA
NM 018974	Homo sapiens unc93 (C.elegans) homolog A (UNC93A), mRNA
NM 014499	Homo sapiens putative purinergic receptor (P2Y10), mRNA
NM 020530	Homo sapiens oncostatin M (OSM), mRNA
NM 020529	Homo sapiens nuclear factor of kappa light polypeptide gene enhancer in B-cells
14141_020323	inhibitor, alpha (NFKBIA), mRNA
NM 014204	Homo sapiens BCL2-related ovarian killer (BOK), mRNA
NM 020527	Homo sapiens HUG1 gene (HUG1), mRNA
NM 006093	Homo sapiens proteoglycan 3 (PRG3), mRNA
NM 020533	Homo sapiens mucolipin 1 (MCOLN1), mRNA
NM 007345	Homo sapiens zinc finger protein 236 (ZNF236), mRNA
NM 002217	Homo sapiens pre-alpha (globulin) inhibitor, H3 polypeptide (ITIH3), mRNA
NM 018693	Homo sapiens vitiligo-associated protein VIT-1 (VIT1), mRNA
NM 006777	Homo sapiens Kaiso (ZNF-kaiso), mRNA
NM 020436	Homo sapiens similar to SALL1 (sal (Drosophila)-like (LOC57167), mRNA
NM 020142	Homo sapiens NADH: ubiquinone oxidoreductase MLRQ subunit homolog
1NIVI_020142	(LOC56901), mRNA
NM 020123	Homo sapiens endomembrane protein emp70 precursor isolog (LOC56889),
14141_020123	mRNA
NM 018845	Homo sapiens stromal cell protein (LOC55974), mRNA
NM 018842	Homo sapiens insulin receptor tyrosine kinase substrate (LOC55971), mRNA
NM 018841	Homo sapiens G-protein gamma-12 subunit (LOC55970), mRNA
NM 018839	Homo sapiens 947 protein (LOC55968), mRNA
NM 016352	Homo sapiens carboxypeptidase A3 (LOC51200), mRNA
NM 016302	Homo sapiens protein x 0001 (LOC51185), mRNA
NM 014332	Homo sapiens small muscle protein, X-linked (SMPX), mRNA
NM 018948	Homo sapiens Gene 33/Mig-6 (MIG-6), mRNA
NM 014587	Homo sapiens SRY (sex determining region Y)-box 8 (SOX8), mRNA
NM 005745	Homo sapiens accessory proteins BAP31/BAP29 (DXS1357E), mRNA
NM 001094	Homo sapiens amiloride-sensitive cation channel 1, neuronal (degenerin)
11111_001094	(ACCN1), mRNA
NM_019609	Homo sapiens metallocarboxypeptidase CPX-1 (CPX-1), mRNA
NM 018844	Homo sapiens B-cell receptor-associated protein BAP29 (BAP29), mRNA
NM 017572	Homo sapiens G protein-coupled receptor kinase 7 (GPRK7), mRNA
NM 016418	Homo sapiens clone FLB5214 (LOC51219), mRNA
NM 016301	Homo sapiens protein x 0004 (LOC51184), mRNA
NM 013387	Homo sapiens ubiquinol-cytochrome c reductase complex (7.2 kD) (HSPC051),
1,11,1 -01,220,	mRNA
NM_020469	Homo sapiens ABO blood group (transferase A, alpha 1-3-N-
14141_020409	acetylgalactosaminyltransferase; transferase B, alpha 1-3-galactosyltransferase)
	(ABO), mRNA
NM 020445	Homo sapiens actin-related protein 3-beta (ARP3BETA), mRNA
NM 020435	Homo sapiens connexin46.6 (CX46.6), mRNA
NM 020426	Homo sapiens lysozyme homolog (LOC57151), mRNA
NM 020379	Homo sapiens 1,2-alpha-mannosidase IC (HMIC), mRNA
NM 020407	Homo sapiens Rh type B glycoprotein (RHBG), mRNA
14141 020407	Tromo sapiens for type D grycoprotein (formo), midak

NM 020406	Homo sapiens polycythemia rubra vera 1; cell surface receptor (PRV1), mRNA
NM 020377	Homo sapiens cysteinyl leukotriene CysLT2 receptor; cDNA PSEC0146 from
	clone PLACE1006979 (LOC57105), mRNA
NM 020355	Homo sapiens HRPAP20 short form (LOC57090), mRNA
NM 020350	Homo sapiens ATRAP protein (ATRAP), mRNA
NM 020380	Homo sapiens AF15q14 protein (AF15Q14), mRNA
NM 020368	Homo sapiens disrupter of silencing 10 (SAS10), mRNA
NM 020344	Homo sapiens solute carrier family 24 (sodium/potassium/calcium exchanger),
_	member 2 (SLC24A2), mRNA
NM 020396	Homo sapiens BCL2-like 10 (apoptosis facilitator) (BCL2L10), mRNA
NM 020384	Homo sapiens claudin 2 (CLDN2), mRNA
NM 007260	Homo sapiens lysophospholipase II (LYPLA2), mRNA
NM 000390	Homo sapiens choroideremia (Rab escort protein 1) (CHM), mRNA
NM 001994	Homo sapiens coagulation factor XIII, B polypeptide (F13B), mRNA
NM 000129	Homo sapiens coagulation factor XIII, A1 polypeptide (F13A1), mRNA
NM 000505	Homo sapiens coagulation factor XII (Hageman factor) (F12), mRNA
NM 000504	Homo sapiens coagulation factor X (F10), mRNA
NM 005509	Homo sapiens Dmx-like 1 (DMXL1), mRNA
NM 001300	Homo sapiens core promoter element binding protein (COPEB), mRNA
NM 012089	Homo sapiens ATP-binding cassette, sub-family B (MDR/TAP), member 10
1411_012005	(ABCB10), nuclear gene encoding mitochondrial protein, mRNA
NM 007188	Homo sapiens ATP-binding cassette, sub-family B (MDR/TAP), member 8
141,1_00,100	(ABCB8), nuclear gene encoding mitochondrial protein, mRNA
NM_005689	Homo sapiens ATP-binding cassette, sub-family B (MDR/TAP), member 6
1111_000000	(ABCB6), nuclear gene encoding mitochondrial protein, mRNA
NM 001216	Homo sapiens carbonic anhydrase IX (CA9), mRNA
NM 000717	Homo sapiens carbonic anhydrase IV (CA4), mRNA
NM 001218	Homo sapiens carbonic anhydrase XII (CA12), mRNA
NM 001217	Homo sapiens carbonic anhydrase XI (CA11), mRNA
NM 006384	Homo sapiens calcium and integrin binding protein (DNA-dependent protein
	kinase interacting protein) (SIP2-28), mRNA
NM 016734	Homo sapiens paired box gene 5 (B-cell lineage specific activator protein)
_	(PAX5), mRNA
NM_000687	Homo sapiens S-adenosylhomocysteine hydrolase (AHCY), mRNA
NM_004482	Homo sapiens UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-
	acetylgalactosaminyltransferase 3 (GalNAc-T3) (GALNT3), mRNA
NM_004481	Homo sapiens UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-
	acetylgalactosaminyltransferase 2 (GalNAc-T2) (GALNT2), mRNA
NM_000512	Homo sapiens galactosamine (N-acetyl)-6-sulfate sulfatase (Morquio syndrome,
	mucopolysaccharidosis type IVA) (GALNS), mRNA
NM_000403	Homo sapiens galactose-4-epimerase, UDP- (GALE), mRNA
NM_020310	Homo sapiens MAX binding protein (MNT), mRNA
NM_006250	Homo sapiens proline-rich protein HaeIII subfamily 1 (PRH1), mRNA
NM_005164	Homo sapiens ATP-binding cassette, sub-family D (ALD), member 2 (ABCD2),
	mRNA
NM_020300	Homo sapiens microsomal glutathione S-transferase 1 (MGST1), mRNA
NM_000728	Homo sapiens calcitonin-related polypeptide, beta (CALCB), mRNA
NM_020127	Homo sapiens tuftelin 1 (TUFT1), mRNA
NM_020040	Homo sapiens tubulin, beta polypeptide 4, member Q (TUBB4Q), mRNA
NM_020126	Homo sapiens sphingosine kinase type 2 isoform (SPHK2), mRNA
NM_020203	Homo sapiens matrix, extracellular phosphoglycoprotein with ASARM motif
1 -	(bone) (MEPE), mRNA

ND ( 020221	Homo sapiens x 010 protein (MDS010), mRNA
NM_020231	Homo sapiens x 010 protein (MDS010), file 14
NM_020132	Homo sapiens lysophosphatidic acid acyltransferase-gamma1 (LPAAT-
	gamma1), mRNA
NM_020246	Homo sapiens cation-chloride cotransporter-interacting protein (LOC56996),
	mRNA
NM_020243	Homo sapiens mitochondrial import receptor Tom22 (LOC56993), mRNA
NM_020240	Homo sapiens non-kinase Cdc42 effector protein SPEC2 (LOC56990), mRNA
NM_020184	Homo sapiens ancient conserved domain protein 4 (LOC56939), mRNA
NM_020178	Homo sapiens Carbonic anhydrase-related protein 10 (LOC56934), mRNA
NM_020155	Homo sapiens chromosome 11 hypothetical protein ORF4 (LOC56834), mRNA
NM 020179	Homo sapiens FN5 protein (FN5), mRNA
NM 020187	Homo sapiens DC12 protein (DC12), mRNA
NM 020156	Homo sapiens core1 UDP-galactose:N-acetylgalactosamine-alpha-R beta 1,3-
_	galactosyltransferase (C1GALT1), mRNA
NM_000352	Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 8
_	(ABCC8), mRNA
NM 000374	Homo sapiens uroporphyrinogen decarboxylase (UROD), mRNA
NM 002872	Homo sapiens ras-related C3 botulinum toxin substrate 2 (rho family, small GTP
	binding protein Rac2) (RAC2), mRNA
NM 004152	Homo sapiens ornithine decarboxylase antizyme 1 (OAZ1), mRNA
NM 002527	Homo sapiens neurotrophin 3 (NTF3), mRNA
NM 002295	Homo sapiens laminin receptor 1 (67kD, ribosomal protein SA) (LAMR1),
1111_002255	mRNA
NM 002293	Homo sapiens laminin, gamma 1 (formerly LAMB2) (LAMC1), mRNA
NM 002292	Homo sapiens laminin, beta 2 (laminin S) (LAMB2), mRNA
NM 002290	Homo sapiens laminin, alpha 4 (LAMA4), mRNA
NM 006192	Homo sapiens paired box gene 1 (PAX1), mRNA
NM 019896	Homo sapiens DNA polymerase epsilon p12 subunit (P12), mRNA
NM_000583	Homo sapiens group-specific component (vitamin D binding protein) (GC),
1111_00000	mRNA
NM_019891	Homo sapiens endoplasmic reticulum oxidoreductin 1-Lbeta (ERO1-L(BETA)),
	mRNA
NM_006705	Homo sapiens growth arrest and DNA-damage-inducible, gamma (GADD45G),
	mRNA
NM 001924	Homo sapiens growth arrest and DNA-damage-inducible, alpha (GADD45A),
	mRNA
NM 019844	Homo sapiens solute carrier family 21 (organic anion transporter), member 8
_======================================	(SLC21A8), mRNA
NM 019644	Homo sapiens testis-specific ankyrin motif containing protein (LOC56311),
	mRNA
NM 019842	Homo sapiens potassium voltage-gated channel, KQT-like subfamily, member 5
	(KCNO5), mRNA
NM_012281	Homo sapiens potassium voltage-gated channel, Shal-related subfamily, member
112.2_012.01	2 (KCND2), mRNA
NM 019857	Homo sapiens CTP synthase II (CTPS2), mRNA
NM 019839	Homo sapiens seven transmembrane receptor BLTR2; leukotriene B4 receptor
1.2.2_013033	BLT2 (BLTR2), mRNA
NM 005757	Homo sapiens C3H-type zinc finger protein; similar to D. melanogaster
1111_000/0/	muscleblind B protein (MBLL), mRNA
NM 004299	Homo sapiens ATP-binding cassette, sub-family B (MDR/TAP), member 7
1111_00 1277	(ABCB7), nuclear gene encoding mitochondrial protein, mRNA
NM 004683	Homo sapiens regucalcin (senescence marker protein-30) (RGN), mRNA
1414 00-1003	1 1101110 Sapieno l'ognosient (Sellescellee market protein-50) (1011); mat 11

	A (OT ATTA) DELA
NM_019618	Homo sapiens interleukin-1 homolog 1 (IL-1H1), mRNA
NM_018950	Homo sapiens major histocompatibility complex, class I, F (HLA-F), mRNA
NM_019610	Homo sapiens hypothetical protein 669 (LOC56267), mRNA
NM_000523	Homo sapiens homeo box D13 (HOXD13), mRNA
NM 019607	Homo sapiens hypothetical protein FLJ11267 (FLJ11267), mRNA
NM 019604	Homo sapiens class-I MHC-restricted T cell associated molecule (CRTAM),
	mRNA
NM 012328	Homo sapiens microvascular endothelial differentiation gene 1 (MDG1), mRNA
NM 013303	Homo sapiens fetal hypothetical protein (HSU84971), mRNA
NM 013298	Homo sapiens hypothetical protein (HSU79252), mRNA
NM 013386	Homo sapiens hypothetical protein (DKFZp586G0123), mRNA
NM 013313	Homo sapiens hypothetical protein (AF060862), mRNA
NM 019116	Homo sapiens similar to ubiquitin binding protein (UBPH), mRNA
NM 018961	Homo sapiens ubiquitin associated and SH3 domain containing, A (UBASH3A),
14141_010501	mRNA
NM 018968	Homo sapiens syntrophin, gamma 2 (SNTG2), mRNA
NM 018967	Homo sapiens syntrophin, gamma 1 (SNTG1), mRNA
NM 018969	Homo sapiens super conserved receptor expressed in brain 3 (SREB3), mRNA
NM 018964	Homo sapiens solute carrier family 37 (glycerol-3-phosphate transporter),
14147 0 1 0 2 0 4	member 1 (SLC37A1), mRNA
NM 018945	Homo sapiens phosphodiesterase 7B (PDE7B), mRNA
NM 019066	Homo sapiens MAGE-like 2 (MAGEL2), mRNA
NM 019060	Homo sapiens NICE-1 protein (NICE-1), mRNA
	Homo sapiens hypothetical protein (LOC55924), mRNA
NM_019099	Homo sapiens spindlin-like (LOC54466), mRNA
NM_019003	Homo sapiens homeo box B6 (HOXB6), mRNA
NM_018952 NM_018951	Homo sapiens homeo box A10 (HOXA10), mRNA
NM 018942	Homo sapiens homeo box (H6 family) 1 (HMX1), mRNA
NM 019109	Homo sapiens beta-1,4 mannosyltransferase (HMT-1), mRNA
NM 019052	Homo sapiens HCR (a-helix coiled-coil rod homologue) (HCR), mRNA
NM 018985	Homo sapiens hypothetical protein (HCGIV.9), mRNA
	Homo sapiens GTP binding protein 2 (GTPBP2), mRNA
NM_019096	Homo sapiens G protein-coupled receptor 14 (GPR14), mRNA
NM_018949	Homo sapiens de protein-coupled receptor 14 (GFR14), micror  Homo sapiens hypothetical protein (FLJ20752), mRNA
NM_019048	Homo sapiens hypothetical protein (FLJ20732), indvA  Homo sapiens hypothetical protein FLJ20674 (FLJ20674), mRNA
NM_019086	Homo sapiens hypothetical protein FLJ20074 (FLJ20074), mRNA
NM_019040	Homo sapiens hypothetical protein (FLJ20498), mRNA
NM_018988	Homo sapiens hypothetical protein (FLJ20330), mRNA
NM_019005	Homo sapiens hypothetical protein (FLJ20323), mRNA
NM_019027	Homo sapiens hypothetical protein (FLJ20273), mRNA
NM_019008	Homo sapiens hypothetical protein (FLJ20232), mRNA
NM_019000	Homo sapiens hypothetical protein (FLJ20152), mRNA
NM_019087	Homo sapiens hypothetical protein FLJ20051 (FLJ20051), mRNA
NM_018996	Homo sapiens hypothetical protein (FLJ20015), mRNA
NM_019021	Homo sapiens hypothetical protein (FLJ20010), mRNA
NM_019018	Homo sapiens hypothetical protein (FLJ11127), mRNA
NM_019084	Homo sapiens hypothetical protein FLJ10895 (FLJ10895), mRNA
NM_019070	Homo sapiens hypothetical protein (FLJ10432), mRNA
NM_019088	Homo sapiens hypothetical protein F23149_1 (F23149_1), mRNA
NM_019002	Homo sapiens ETAA16 protein (ETAA16), mRNA
NM_019114	Homo sapiens EHM2 gene (EHM2), mRNA
NM_018973	Homo sapiens dolichyl-phosphate mannosyltransferase polypeptide 3 (DPM3),
-	mRNA

	The state of the s
	Homo sapiens DAZ associated protein 1 (DAZAP1), mRNA
	Homo sapiens cyclic nucleotide gated channel beta 3 (CNGB3), mRNA
NM_018958	Homo sapiens chromosome 15 open reading frame 2 (C15ORF2), mRNA
NM_000379	Homo sapiens xanthene dehydrogenase (XDH), mRNA
NM 000552	Homo sapiens von Willebrand factor (VWF), mRNA
NM 000362	Homo sapiens tissue inhibitor of metalloproteinase 3 (Sorsby fundus dystrophy,
_	pseudoinflammatory) (TIMP3), mRNA
NM 003255	Homo sapiens tissue inhibitor of metalloproteinase 2 (TIMP2), mRNA
NM_003001	Homo sapiens succinate dehydrogenase complex, subunit C, integral membrane protein, 15kD (SDHC), nuclear gene encoding mitochondrial protein, mRNA
NM_003000	Homo sapiens succinate dehydrogenase complex, subunit B, iron sulfur (Ip) (SDHB), nuclear gene encoding mitochondrial protein, mRNA
NM 006745	Homo sapiens sterol-C4-methyl oxidase-like (SC4MOL), mRNA
	Homo sapiens putative GTP-binding protein similar to RAY/RAB1C (RAYL),
NM_006860	mRNA
NM_000531	Homo sapiens ornithine carbamoyltransferase (OTC), nuclear gene encoding mitochondrial protein, mRNA
NM 000607	Homo sapiens orosomucoid 1 (ORM1), mRNA
NM 002538	Homo sapiens occludin (OCLN), mRNA
NM 002301	Homo sapiens lactate dehydrogenase C (LDHC), transcript variant 1, mRNA
NM 017448	Homo sapiens lactate dehydrogenase C (LDHC), transcript variant 2, mRNA
NM 000892	Homo sapiens kallikrein B, plasma (Fletcher factor) 1 (KLKB1), mRNA
NM 002193	Homo sapiens inhibin, beta B (activin AB beta polypeptide) (INHBB), mRNA
NM 002191	Homo sapiens inhibin, alpha (INHA), mRNA
NM 002015	Homo sapiens forkhead box O1A (rhabdomyosarcoma) (FOXO1A), mRNA
NM 004473	Homo sapiens forkhead box E1 (thyroid transcription factor 2) (FOXE1), mRNA
NM 000804	Homo sapiens folate receptor 3 (gamma) (FOLR3), mRNA
NM 000803	Homo sapiens folate receptor 2 (fetal) (FOLR2), mRNA
NM 004742	Homo sapiens BAI1-associated protein 1 (BAIAP1), mRNA
NM 004925	Homo saniens aquaporin 3 (AOP3), mRNA
NM_007182	Homo sapiens Ras association (RalGDS/AF-6) domain family 1 (RASSF1), mRNA
NM_018941	Homo sapiens ceroid-lipofuscinosis, neuronal 8 (epilepsy, progressive with
NWI_010941	mental retardation) (CLN8), mRNA
NM 016936	Homo sapiens ubinuclein 1 (UBN1), mRNA
NM 012406	Homo sapiens PR domain containing 4 (PRDM4), mRNA
NM 018728	Homo sapiens myosin 5C (MYO5C), mRNA
NM_017540	Homo sapiens hypothetical protein DKFZp586H0623 (DKFZp586H0623),
NM_018651	mRNA Homo sapiens zinc finger protein (ZFP), mRNA
NM_017503	Homo sapiens surfeit 2 (SURF2), mRNA
NM_018419	Homo sapiens SRY (sex determining region Y)-box 18 (SOX18), mRNA
NM_018427	Homo sapiens RNA polymerase I transcription factor RRN3 (RRN3), mRNA
NM_018545	Homo sapiens hypothetical protein PRO2955 (PRO2955), mRNA
NM_018525	Homo sapiens hypothetical protein PRO2369 (PRO2369), mRNA
NM 018520	Homo sapiens hypothetical protein PRO2268 (PRO2268), mRNA
NM 018605	Homo sapiens hypothetical protein PRO1777 (PRO1777), mRNA
NM 018573	Homo sapiens hypothetical protein PRO1068 (PRO1068), mRNA
	Homo sapiens hypothetical protein PRO1051 (PRO1051), mRNA
NM_018572	Homo sapiens hypothetical protein PRO1051 (PRO1051), mRNA
	Homo sapiens hypothetical protein PRO1051 (PRO1051), mRNA Homo sapiens hypothetical protein PRO0971 (PRO0971), mRNA Homo sapiens hypothetical protein PRO0800 (PRO0800), mRNA Homo sapiens hypothetical protein PRO0758 (PRO0758), mRNA

NM 018699	Homo sapiens PR domain containing 5 (PRDM5), mRNA
NM 017534	Homo sapiens myosin, heavy polypeptide 2, skeletal muscle, adult (MYH2),
14141-011224	mRNA
NM_018461	Homo sapiens uncharacterized hematopoietic stem/progenitor cells protein
11111_010401	MDS026 (MDS026), mRNA
NM 018559	Homo sapiens lipopolysaccharide specific response-7 protein (LSR7), mRNA
NM 018694	Homo sapiens HSVI binding protein (LOC55913), mRNA
NM 018663	Homo sapiens 22kDa peroxisomal membrane protein-like (LOC55895), mRNA
NM 018640	Homo sapiens neuronal specific transcription factor DAT1 (LOC55885), mRNA
NM 018639	Homo sapiens CS box-containing WD protein (LOC55884), mRNA
NM 018449	Homo sapiens AD-012 protein (LOC55833), mRNA
NM 018658	Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 16
1414_010050	(KCNJ16), mRNA
NM_018671	Homo sapiens hypothetical protein (IRO039700), mRNA
NM 018439	Homo sapiens hypothetical protein IMPACT (IMPACT), mRNA
NM 017521	Homo sapiens FEV protein (HSRNAFEV), mRNA
NM 017526	Homo sapiens leptin receptor gene-related protein (HSOBRGRP), mRNA
NM 017513	Homo sapiens metaphase chromosome protein 1 (HSMCR30), mRNA
NM 017532	Homo sapiens p65 protein (HSAJ2425), mRNA
NM 018682	Homo sapiens hypothetical protein HDCMC04P (HDCMC04P), mRNA
NM 018680	Homo sapiens hypothetical protein HDCGC21P (HDCGC21P), mRNA
NM 018428	Homo sapiens hepatocellular carcinoma-associated antigen 66 (HCA66), mRNA
NM 017528	Homo sapiens putative methyltransferase (HASJ4442), mRNA
NM 017964	Homo sapiens hypothetical protein FLJ20837 (FLJ20837), mRNA
NM 017952	Homo sapiens hypothetical protein FLJ20758 (FLJ20758), mRNA
NM 017936	Homo sapiens hypothetical protein FLJ20707 (FLJ20707), mRNA
NM 017933	Homo sapiens hypothetical protein FLJ20701 (FLJ20701), mRNA
NM 017931	Homo sapiens hypothetical protein FLJ20699 (FLJ20699), mRNA
NM 017911	Homo sapiens hypothetical protein FLJ20635 (FLJ20635), mRNA
NM 017898	Homo sapiens hypothetical protein FLJ20605 (FLJ20605), mRNA
NM 017888	Homo sapiens hypothetical protein FLJ20581 (FLJ20581), mRNA
NM 017865	Homo sapiens hypothetical protein FLJ20531 (FLJ20531), mRNA
NM 017855	Homo sapiens hypothetical protein FLJ20513 (FLJ20513), mRNA
NM_017849	Homo sapiens hypothetical protein FLJ20507 (FLJ20507), mRNA
NM_017845	Homo sapiens hypothetical protein FLJ20502 (FLJ20502), mRNA
NM_017842	Homo sapiens hypothetical protein FLJ20489 (FLJ20489), mRNA
NM_017820	Homo sapiens hypothetical protein FLJ20433 (FLJ20433), mRNA
NM_017806	Homo sapiens hypothetical protein FLJ20406 (FLJ20406), mRNA
NM_017800	Homo sapiens hypothetical protein FLJ20393 (FLJ20393), mRNA
NM_017795	Homo sapiens hypothetical protein FLJ20378 (FLJ20378), mRNA
NM_017794	Homo sapiens hypothetical protein FLJ20375 (FLJ20375), mRNA
NM_017768	Homo sapiens hypothetical protein FLJ20331 (FLJ20331), mRNA
NM_017757	Homo sapiens hypothetical protein FLJ20307 (FLJ20307), mRNA
NM_017749	Homo sapiens hypothetical protein FLJ20294 (FLJ20294), mRNA
NM_017733	Homo sapiens hypothetical protein FLJ20265 (FLJ20265), mRNA
NM_017732	Homo sapiens hypothetical protein FLJ20262 (FLJ20262), mRNA
NM_017730	Homo sapiens hypothetical protein FLJ20259 (FLJ20259), mRNA
NM_017723	Homo sapiens hypothetical protein FLJ20245 (FLJ20245), mRNA
NM_017720	Homo sapiens hypothetical protein FLJ20234 (FLJ20234), mRNA
NM_017715	Homo sapiens hypothetical protein FLJ20216 (FLJ20216), mRNA
NM_017667	Homo sapiens hypothetical protein FLJ20097 (FLJ20097), mRNA
NM 017652	Homo sapiens hypothetical protein FLJ20070 (FLJ20070), mRNA

	CT YOUR DAY
NM_017635	Homo sapiens hypothetical protein FLJ20039 (FLJ20039), mRNA
NM_017632	Homo sapiens hypothetical protein FLJ20036 (FLJ20036), mRNA
NM_017624	Homo sapiens hypothetical protein FLJ20019 (FLJ20019), mRNA
NM_017623	Homo sapiens hypothetical protein FLJ20018 (FLJ20018), mRNA
NM_018390	Homo sapiens hypothetical protein FLJ11323 (FLJ11323), mRNA
NM 018382	Homo sapiens hypothetical protein FLJ11292 (FLJ11292), mRNA
NM 018337	Homo sapiens hypothetical protein FLJ11137 (FLJ11137), mRNA
NM 018320	Homo sapiens hypothetical protein FLJ11099 (FLJ11099), mRNA
NM 018317	Homo sapiens hypothetical protein FLJ11082 (FLJ11082), mRNA
NM 018301	Homo sapiens hypothetical protein FLJ11016 (FLJ11016), mRNA
NM 018295	Homo sapiens hypothetical protein FLJ11000 (FLJ11000), mRNA
NM 018291	Homo sapiens hypothetical protein FLJ10986 (FLJ10986), mRNA
NM 018290	Homo sapiens hypothetical protein FLJ10983 (FLJ10983), mRNA
NM 018280	Homo sapiens hypothetical protein FLJ10945 (FLJ10945), mRNA
NM 018266	Homo sapiens hypothetical protein FLJ10902 (FLJ10902), mRNA
NM 018263	Homo sapiens hypothetical protein FLJ10898 (FLJ10898), mRNA
NM 018249	Homo sapiens hypothetical protein FLJ10867 (FLJ10867), mRNA
NM 018233	Homo sapiens hypothetical protein FLJ10826 (FLJ10826), mRNA
NM 018202	Homo sapiens hypothetical protein FLJ10747 (FLJ10747), mRNA
NM 018194	Homo sapiens hypothetical protein FLJ10724 (FLJ10724), mRNA
NM 018191	Homo sapiens hypothetical protein FLJ10716 (FLJ10716), mRNA
NM 018134	Homo sapiens hypothetical protein FLJ10547 (FLJ10547), mRNA
NM 018131	Homo sapiens hypothetical protein FLJ10540 (FLJ10540), mRNA
NM 018124	Homo sapiens hypothetical protein FLJ10520 (FLJ10520), mRNA
NM 018114	Homo sapiens hypothetical protein FLJ10496 (FLJ10496), mRNA
NM 018107	Homo sapiens hypothetical protein FLJ10482 (FLJ10482), mRNA
NM 018098	Homo sapiens hypothetical protein FLJ10461 (FLJ10461), mRNA
NM 018085	Homo sapiens hypothetical protein FLJ10402 (FLJ10402), mRNA
NM 018079	Homo sapiens hypothetical protein FLJ10379 (FLJ10379), mRNA
NM 018063	Homo sapiens hypothetical protein FLJ10339 (FLJ10339), mRNA
NM 018062	Homo sapiens hypothetical protein FLJ10335 (FLJ10335), mRNA
NM 018059	Homo sapiens hypothetical protein FLJ10324 (FLJ10324), mRNA
NM 018053	Homo sapiens hypothetical protein FLJ10307 (FLJ10307), mRNA
NM 018046	Homo sapiens hypothetical protein FLJ10283 (FLJ10283), mRNA
NM 018006	Homo sapiens hypothetical protein FLJ10140 (FLJ10140), mRNA
NM_018004	Homo sapiens hypothetical protein FLJ10134 (FLJ10134), mRNA
NM 017999	Homo sapiens hypothetical protein FLJ10111 (FLJ10111), mRNA
NM 017992	Homo sapiens hypothetical protein FLJ10083 (FLJ10083), mRNA
NM 017991	Homo sapiens hypothetical protein FLJ10081 (FLJ10081), mRNA
NM 017979	Homo sapiens hypothetical protein FLJ10043 (FLJ10043), mRNA
NM 017975	Homo sapiens hypothetical protein FLJ10036 (FLJ10036), mRNA
NM 017973	Homo sapiens hypothetical protein FLJ10034 (FLJ10034), mRNA
NM 017610	Homo sapiens hypothetical protein DKFZp761D081 (DKFZp761D081), mRNA
NM 018457	Homo sapiens DKFZp564J157 protein (DKFZP564J157), mRNA
NM 017590	Homo sapiens hypothetical protein DKFZp434K0920 (DKFZp434K0920),
14141_01/390	mRNA
NM 017566	Homo sapiens hypothetical protein DKFZp434G0522 (DKFZp434G0522),
14141_017500	mRNA
NM_017612	Homo sapiens hypothetical protein DKFZp434E2220 (DKFZp434E2220),
14141_017012	mRNA
NM 018641	Homo sapiens chondroitin 4-O-sulfotransferase 2 (C4S-2), mRNA
NM 018659	Homo sapiens cytokine-like protein C17 (C17), mRNA
11171 010000	Total and a second seco

NM 018656	Homo sapiens bladder cancer overexpressed protein (BLOV1), mRNA
NM 018702	Homo sapiens double-stranded RNA specific adenosine deaminase (ADAR3),
1NM_018702	mRNA
NM 014160	Homo sapiens HSPC070 protein (HSPC070), mRNA
NM_004288	Homo sapiens pleckstrin homology, Sec7 and coiled/coil domains, binding
14141_00-1200	protein (PSCDBP), mRNA
NM 004060	Homo sapiens cyclin G1 (CCNG1), mRNA
NM 006521	Homo sapiens transcription factor binding to IGHM enhancer 3 (TFE3), mRNA
NM 007035	Homo sapiens keratocan (KERA), mRNA
NM 000546	Homo sapiens tumor protein p53 (Li-Fraumeni syndrome) (TP53), mRNA
NM 003015	Homo sapiens secreted frizzled-related protein 5 (SFRP5), mRNA
	Homo sapiens secreted frizzled-related protein 1 (SFRP1), mRNA
NM_003012	Homo sapiens ubiquitin specific protease 18 (USP18), mRNA
NM_017414	Homo sapiens ubiquitin associated protein (UBAP), mRNA
NM_016525	Homo sapiens toll-like receptor 9 (TLR9), mRNA
NM_017442	Homo sapiens ton-like receptor y (TERO), interview Homo sapiens polymerase (DNA directed), alpha (POLA), mRNA
NM_016937	Homo sapiens NADPH oxidase 4 (NOX4), mRNA
NM_016931	Homo sapiens NADPH Oxidase 4 (NOX4), intervi
NM_017433	Homo sapiens myosin IIIA (MYO3A), mRNA Homo sapiens junctional adhesion molecule (JAM), mRNA
NM_016946	Homo sapiens junctional adnessor molecule (JAW), mixty
NM_005536	Homo sapiens inositol(myo)-1(or 4)-monophosphatase 1 (IMPA1), mRNA
NM_017410	Homo sapiens homeo box C13 (HOXC13), mRNA
NM_017409	Homo sapiens homeo box C10 (HOXC10), mRNA
NM_015922	Homo sapiens NAD(P) dependent steroid dehydrogenase-like; H105e3
	(H105E3), mRNA
NM_004129	Homo sapiens guanylate cyclase 1, soluble, beta 2 (GUCY1B2), mRNA
NM_017423	Homo sapiens UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-
	acetylgalactosaminyltransferase 7 (GalNAc-T7) (GALNT7), mRNA
NM_016947	Homo sapiens G8 protein (G8), mRNA
NM_017434	Homo sapiens dual oxidase 1 (DUOX1), mRNA
NM_012143	Homo sapiens tuftelin-interacting protein (TIP39), mRNA
NM_017418	Homo sapiens deleted in esophageal cancer 1 (DEC1), mRNA
NM_016929	Homo sapiens chloride intracellular channel 5 (CLIC5), mRNA
NM_017413	Homo sapiens apelin; peptide ligand for APJ receptor (APELIN), mRNA
NM_000477	Homo sapiens albumin (ALB), mRNA
NM_007235	Homo sapiens exportin, tRNA (nuclear export receptor for tRNAs) (XPOT),
	mRNA
NM_004585	Homo sapiens retinoic acid receptor responder (tazarotene induced) 3
	(RARRES3), mRNA
NM_002134	Homo sapiens heme oxygenase (decycling) 2 (HMOX2), mRNA
NM_002100	Homo sapiens glycophorin B (includes Ss blood group) (GYPB), mRNA
NM_002099	Homo sapiens glycophorin A (includes MN blood group) (GYPA), mRNA
NM_005708	Homo sapiens glypican 6 (GPC6), mRNA
NM_013280	Homo sapiens fibronectin leucine rich transmembrane protein 1 (FLRT1),
	mRNA P (CDD) DNA
NM_001304	Homo sapiens carboxypeptidase D (CPD), mRNA
NM_013410	Homo sapiens adenylate kinase 3 (AK3), nuclear gene encoding mitochondrial
	protein, mRNA
NM_002161	Homo sapiens isoleucine-tRNA synthetase (IARS), transcript variant short,
	mRNA
NM_013417	Homo sapiens isoleucine-tRNA synthetase (IARS), transcript variant long,
	mRNA
NM 015836	Homo sapiens tryptophanyl tRNA synthetase 2 (mitochondrial) (WARS2),

	DNA
	nuclear gene encoding mitochondrial protein, mRNA
NM_004992	Homo sapiens methyl CpG binding protein 2 (Rett syndrome) (MECP2), mRNA
NM_003926	Homo sapiens methyl-CpG binding domain protein 3 (MBD3), mRNA
NM_006150	Homo sapiens LIM domain only 6 (LMO6), mRNA
NM_013431	Homo sapiens killer cell lectin-like receptor subfamily C, member 4 (KLRC4), mRNA
NM 001427	Homo sapiens engrailed homolog 2 (EN2), mRNA
NM 001426	Homo sapiens engrailed homolog 1 (EN1), mRNA
NM 003445	Homo sapiens zinc finger protein 155 (pHZ-96) (ZNF155), mRNA
NM 016220	Homo sapiens zinc finger protein (ZFD25) (ZFD25), mRNA
NM 015855	Homo sapiens Wilms tumor associated protein (WIT-1), mRNA
NM 015873	Homo sapiens villin-like (VILL), mRNA
NM 016379	Homo sapiens variable charge protein on X with eight repeats (VCX-8r), mRNA
NM 016378	Homo sapiens variable charge protein on X with two repeats (VCX-2r), mRNA
NM 016437	Homo sapiens tubulin, gamma 2 (TUBG2), mRNA
NM 016575	Homo sapiens TU12B1-TY protein (TU12B1-TY), mRNA
NM 016089	Homo sapiens KRAB-zinc finger protein SZF1-1 (SZF1), mRNA
NM_013272	Homo sapiens solute carrier family 21 (organic anion transporter), member 11
NW_013272	(SLC21A11), mRNA
NM 015926	Homo sapiens putative secreted protein (SIG11), mRNA
	Homo sapiens SH3 and PX domain-containing protein SH3PX1 (SH3PX1),
NM_016224	mRNA
NM_016276	Homo sapiens serum/glucocorticoid regulated kinase 2 (SGK2), mRNA
NM_015884	Homo sapiens S2P protein (S2P), mRNA
NM_016356	Homo sapiens RU2S (RU2), mRNA
NM_016321	Homo sapiens Rh type C glycoprotein (RHCG), mRNA
NM_015900	Homo sapiens phosphatidylserine-specific phospholipase A1alpha (PS-PLA1), mRNA
NM 016533	Homo sapiens ninjurin 2 (NINJ2), mRNA
NM 016641	Homo sapiens membrane interacting protein of RGS16 (MIR16), mRNA
NM 014319	Homo sapiens integral inner nuclear membrane protein (MAN1), mRNA
NM_016249	Homo sapiens melanoma antigen, family E, 1, cancer/testis specific (MAGEE1), mRNA
NM 016153	Homo sapiens LW-1 (LW-1), mRNA
NM 016551	Homo sapiens seven transmembrane protein TM7SF3 (TM7SF3), mRNA
NM_016529	Homo sapiens ATPase, aminophospholipid transporter-like, Class I, type 8A, member 2 (ATP8A2), mRNA
ND4 016422	Homo sapiens synoretin (LOC51749), mRNA
NM_016432	Homo sapiens synoretin (LOC51749), mRNA  Homo sapiens ghrelin precursor (LOC51738), mRNA
NM_016362	Homo sapiens grienn precursor (LOC51738), mRNA  Homo sapiens Kruppel-like factor (LOC51713), mRNA
NM_016270	Homo sapiens Kruppei-like factor (LOCS1715), filedya
NM_016243	Homo sapiens cytochrome b5 reductase 1 (B5R.1) (LOC51706), mRNA
NM_016231	Homo sapiens nemo-like kinase (LOC51701), mRNA
NM_016225	Homo sapiens RhD type IIIa protein (LOC51698), mRNA
NM_016219	Homo sapiens alpha 1,2-mannosidase (LOC51697), mRNA
NM_016217	Homo sapiens hHDC for homolog of Drosophila headcase (LOC51696), mRNA
NM_016199	Homo sapiens U6 snRNA-associated Sm-like protein LSm7 (LOC51690), mRNA
NM 016171	Homo sapiens prothymosin a14 (LOC51685), mRNA
1 - 1-1-	Homo sapiens MAGUK protein p55T; Protein Associated with Lins 2
NM_016447	(I OC51678) mRNA
NM_016447 NM_016126	(LOC51678), mRNA  Homo sapiens HSPCO34 protein (LOC51668), mRNA

	CC7.140 (T.O.C51652) mPN/A
NM_016079	Homo sapiens CGI-149 protein (LOC51652), mRNA
NM_016062	Homo sapiens CGI-128 protein (LOC51647), mRNA
NM_016057	Homo sapiens CGI-120 protein (LOC51644), mRNA
NM_016056	Homo sapiens CGI-119 protein (LOC51643), mRNA
NM_016047	Homo sapiens CGI-110 protein (LOC51639), mRNA
NM_016016	Homo sapiens CGI-69 protein (LOC51629), mRNA
NM_016008	Homo sapiens CGI-60 protein (LOC51626), mRNA
NM 015995	Homo sapiens Kruppel-like factor 13 (KLF13), mRNA
NM 015980	Homo sapiens HMP19 protein (LOC51617), mRNA
NM 015958	Homo sapiens CGI-30 protein (LOC51611), mRNA
NM 015941	Homo sapiens CGI-11 protein (LOC51606), mRNA
NM 015937	Homo sapiens CGI-06 protein (LOC51604), mRNA
NM 015929	Homo sapiens lipovltransferase (LOC51601), mRNA
NM 015921	Homo sapiens divalent cation tolerant protein CUTA (LOC51596), mRNA
NM 015908	Homo sapiens arsenate resistance protein ARS2 (ARS2), mRNA
NM 015875	Homo sapiens unnamed HERV-H protein (LOC51581), mRNA
NM 015874	Homo sapiens H-2K binding factor-2 (LOC51580), mRNA
NM 016283	Homo saniens adrenal gland protein AD-004 (LOC51578), mRNA
NM 016644	Homo sapiens mesenchymal stem cell protein DSC54 (LOC51334), mRNA
NM 016643	Homo sapiens mesenchymal stem cell protein DSC43 (LOC51333), mRNA
NM 016642	Homo sapiens beta V spectrin (BSPECV), mRNA
NM 016638	Homo sapiens SRp25 nuclear protein (LOC51329), mRNA
NM 016637	Homo sapiens neaml (LOC51328), mRNA
NM 016633	Homo sapiens EDRF protein (LOC51327), mRNA
NM 016625	Homo sapiens hypothetical protein (LOC51319), mRNA
NM 016622	Homo sapiens hypothetical protein (LOC51318), mRNA
NM 016621	Homo sapiens hypothetical protein (LOC51317), mRNA
NM_016609	Homo sapiens hBOIT for potent brain type organic ion transporter (LOC51310),
11112_01000	mRNA
NM 016606	Homo sapiens SGC32445 protein (LOC51308), mRNA
NM_016591	Homo sapiens core 2 beta-1,6-N-acetylglucosaminyltransferase 3 (LOC51301),
	mRNA
NM 016585	Homo sapiens testicular haploid expressed gene (THEG), mRNA
NM 016573	Homo saniens Gem-interacting protein (LOC51291), mRNA
NM_016568	Homo sapiens G-protein coupled receptor SALPR; somatostatin and angiotensin-
	like peptide receptor (LOC51289), mRNA
NM 016566	Homo sapiens pparl (LOC51288), mRNA
NM 016563	Homo sapiens Ris (LOC51285), mRNA
NM 016548	Homo sapiens golgi membrane protein GP73 (LOC51280), mRNA
NM 016499	Homo sapiens hypothetical protein (LOC51259), mRNA
NM 016490	Homo sapiens hypothetical protein (LOC51252), mRNA
NM 016466	Homo sapiens hypothetical protein (LOC51239), mRNA
NM 016459	Homo sapiens hypothetical protein (LOC51237), mRNA
NM_016449	Homo sapiens hypothetical protein (LOC51233), mRNA
NM 016440	Homo sapiens VRK3 for vaccinia related kinase 3 (LOC51231), mRNA
NM 016427	Homo sapiens transcription elongation factor (SIII) elongin A2 (TCEB3L),
	mRNA
NM 016423	Homo saniens zinc finger protein 219 (ZNF219), mRNA
NM 016361	Homo sapiens LPAP for lysophosphatidic acid phosphatase (LOC51205),
	mRNA
NM 016353	Homo sapiens rec (LOC51201), mRNA
NM 016349	Homo sapiens susceptibility protein NSG-x (LOC51198), mRNA

	0.0051100 mPNA
NM_016341	Homo sapiens pancreas-enriched phospholipase C (LOC51196), mRNA
NM_016323	Homo sapiens cyclin-E binding protein 1 (LOC51191), mRNA
NM_016317	Homo sapiens neutral sphingomyelinase (LOC51190), mRNA
NM_016286	Homo sapiens carbonyl reductase (LOC51181), mRNA
NM_016269	Homo sapiens lymphoid enhancer binding factor-1 (LOC51176), mRNA
NM_016245	Homo sapiens retinal short-chain dehydrogenase/reductase retSDR2
	(LOC51170), mRNA
NM_016241	Homo sapiens endomucin-1 (LOC51169), mRNA
NM_016230	Homo sapiens flavohemoprotein b5+b5R (LOC51167), mRNA
NM_016221	Homo sapiens dynactin p62 subunit (LOC51164), mRNA
NM_016215	Homo sapiens NEU1 protein (LOC51162), mRNA
NM_016210	Homo sapiens g20 protein (LOC51161), mRNA
NM_016161	Homo sapiens alpha-1,4-N-acetylglucosaminyltransferase (LOC51146), mRNA
NM_016123	Homo sapiens putative protein kinase NY-REN-64 antigen (LOC51135), mRNA
NM_016120	Homo sapiens putative ring zinc finger protein NY-REN-43 antigen
_	(LOC51132), mRNA
NM_016033	Homo sapiens CGI-90 protein (LOC51115), mRNA
NM_016032	Homo sapiens CGI-89 protein (LOC51114), mRNA
NM 016030.	Homo sapiens CGI-87 protein (LOC51112), mRNA
NM 016028	Homo sapiens CGI-85 protein (LOC51111), mRNA
NM 016027	Homo sapiens CGI-83 protein (LOC51110), mRNA
NM 016022	Homo sapiens CGI-78 protein (LOC51107), mRNA
NM 016018	Homo sapiens CGI-72 protein (LOC51105), mRNA
NM 016013	Homo sapiens CGI-65 protein (LOC51103), mRNA
NM 016011	Homo sapiens CGI-63 protein (LOC51102), mRNA
NM 016006	Homo sapiens CGI-58 protein (LOC51099), mRNA
NM 015999	Homo sapiens CGI-45 protein (LOC51094), mRNA
NM 015982	Homo sapiens germ cell specific Y-box binding protein (LOC51087), mRNA
NM 015963	Homo sapiens CGI-36 protein (LOC51078), mRNA
NM 015959	Homo sapiens CGI-31 protein (LOC51075), mRNA
NM 015950	Homo sapiens CGI-22 protein (LOC51069), mRNA
NM 015938	Homo sapiens CGI-07 protein (LOC51068), mRNA
NM 015916	Homo sapiens hypothetical protein (LOC51063), mRNA
NM 015914	Homo sapiens hypothetical protein (LOC51061), mRNA
NM 015910	Homo sapiens hypothetical protein (LOC51057), mRNA
NM 015901	Homo sapiens unknown (LOC51055), mRNA
NM 015893	Homo sapiens preproprolactin-releasing peptide (LOC51052), mRNA
NM 015887	Homo sapiens putative peroxisome microbody protein 175.1 (LOC51051),
	mRNA
NM 015880	Homo sapiens RIG-like 14-1 (LOC51047), mRNA
NM 015877	Homo sapiens Kruppel-associated box protein (LOC51045), mRNA
NM 015863	Homo sapiens surfactant protein B (LOC51041), mRNA
NM 015854	Homo sapiens retinoic acid receptor-beta associated open reading frame
	(LOC51036), mRNA
NM 015849	Homo sapiens pancreatic elastase IIB (LOC51032), mRNA
NM 016075	Homo sapiens CGI-145 protein (LOC51028), mRNA
NM 016074	Homo sapiens CGI-143 protein (LOC51027), mRNA
NM 016063	Homo sapiens CGI-130 protein (LOC51020), mRNA
NM_016048	Homo sapiens CGI-111 protein (LOC51015), mRNA
NM 016044	Homo sapiens CGI-105 protein (LOC51011), mRNA
NM 015947	Homo sapiens CGI-18 protein (LOC51008), mRNA
NM 016058	Homo sapiens CGI-121 protein (LOC51002), mRNA
1111 010000	

NM 015948	Homo sapiens CGI-19 protein (LOC51000), mRNA
NM 016040	Homo sapiens CGI-100 protein (LOC50999), mRNA
NM 016571	Homo sapiens lengsin (LGS), mRNA
NM 015868	Homo sapiens NK-receptor (KIR-023GB), mRNA
NM 016281	Homo sapiens STE20-like kinase (JIK), mRNA
NM_016358	Homo seniens iroquois homeobox protein 4 (IRX4), mRNA
NM 016291	Homo sapiens mammalian inositol hexakisphosphate kinase 2 (IPOK2), IIKNA
NM 015848	Homo sapiens cytokeratin 2 (HUMCYT2A), mRNA
NM 016506	Homo sapiens hypothetical protein (HSPC252), mRNA
NM 016498	Homo sapiens hypothetical protein (HSPC242), mRNA
NM 016460	Homo sapiens hypothetical protein (HSPC192), mRNA
NM 016390	Homo sapiens hypothetical protein (HSPC109), mRNA
NM 016091	Homo sapiens HSPC025 (HSPC025), mRNA
NM_016522	Homo saniens neurotrimin (HNT), mRNA
NM 016258	Homo sapiens high-glucose-regulated protein 8 (HGRG8), mRNA
NM 016173	Homo sapiens HEMK homolog 7kb (HEMK), mRNA
NM 016516	Homo sapiens tumor antigen SLP-8p (HCC8), mRNA
NM 016540	Homo sapiens G protein-coupled receptor 72 (GPR72), mRNA
NM 012196	Home capiens G antigen 8 (GAGE8), mRNA
NM 015898	Homo saniens HIV-1 inducer of short transcripts binding protein (FBI1), mRNA
NM 016357	Homo sapiens epithelial protein lost in neoplasm beta (EPLIN), mRNA
NM 016218	Homo sapiens polymerase (DNA-directed) kappa (POLK), mRNA
NM 016240	Homo sapiens CSR1 protein (CSR1), mRNA
NM 016073	Homo sapiens CGI-142 (CGI-142), mRNA
NM 016315	Homo sapiens CED-6 protein (CED-6), mRNA
NM 016620	Homo sapiens hypothetical protein (BM-005), mRNA
NM 015896	Homo saniens BLu protein (BLu), mRNA
NM 016426	Homo ganiens G-2 and S-phase expressed 1 (GTSE1), mRNA
NM_015928	Homo sapiens androgen-induced prostate proliferative shutoff associated protein
TVIVI_013528	(AS3) mRNA
NM 016238	Homo sapiens anaphase-promoting complex subunit 7 (APC7), mRNA
NM 016376	Homo sapiens ANKHZN protein (ANKHZN), mRNA
NM 016282	Homo sapiens adenylate kinase 3 alpha like (AKL3L), mRNA
NM 016453	Homo sapiens SH3 protein (AF3P21), mRNA
NM 016614	Homo saniens TRAF and TNF receptor-associated protein (AD022), mRNA
NM_015365	Homo sapiens Alport syndrome, mental retardation, midface hypoplasia and
14141_015505	elliptocytosis chromosomal region, gene 1 (AMMECR1), mRNA
NM 007126	Homo sapiens valosin-containing protein (VCP), mRNA
NM 001059	Homo saniens tachykinin recentor 3 (TACR3), mRNA
NM 005963	Homo sapiens myosin, heavy polypeptide 1, skeletal muscle, adult (MYH1),
14141_0005505	mRNA
NM 005561	Homo sapiens lysosomal-associated membrane protein 1 (LAMP1), mRNA
NM 006407	Homo saniens vitamin A responsive; cytoskeleton related (JWA), mRNA
NM 000407	Homo seniens glutathione S-transferase theta 2 (GST12), micha
NM 002046	Homo saniens glyceraldehyde-3-phosphate dehydrogenase (GAPD), mRNA
NM 001953	Homo sapiens endothelial cell growth factor 1 (platelet-derived) (ECGF1),
141AT_001322	mDNIA
NIM 000027	Homo sapiens ATP-binding cassette, sub-family B (MDR/TAP), member 1
NM_000927	(ABCB1), mRNA
NM 015686	Homo sapiens TED protein (TED), mRNA
	Homo sapiens STG protein (STG), mRNA
NM_014070 NM_014069	Homo sapiens SPR1 protein (SPR1), mRNA
14141_014009	Home supram or an proven (or)

ND4 014069	Homo sapiens SEEK1 protein (SEEK1), mRNA
NM_014068	Homo sapiens PTD011 protein (PTD011), mRNA
NM_014051	
NM_014109	Homo sapiens PRO2000 protein (PRO2000), mRNA  Homo sapiens PRO1992 protein (PRO1992), mRNA
NM_014107	
NM_014095	Homo sapiens PRO1600 protein (PRO1600), mRNA
NM_014084	Homo sapiens PRO0806 protein (PRO0806), mRNA
NM_014130	Homo sapiens PRO0483 protein (PRO0483), mRNA
NM_014082	Homo sapiens PRO0397 protein (PRO0397), mRNA
NM_014125	Homo sapiens PRO0327 protein (PRO0327), mRNA
NM_014081	Homo sapiens PRO0297 protein (PRO0297), mRNA
NM_014037	Homo sapiens NTT5 protein (NTT5), mRNA  Homo sapiens MIL1 protein (MIL1), nuclear gene encoding mitochondrial
NM_015367	protein, mRNA
NM 014060	Homo sapiens MCT-1 protein (MCT-1), mRNA
NM 014892	Homo sapiens KIAA1116 protein (KIAA1116), mRNA
NM 014968	Homo sapiens KIAA1104 protein (KIAA1104), mRNA
NM 014915	Homo sapiens KIAA1104 protein (KIAA1104), mRNA
NM 014911	Homo sapiens KIAA1048 protein (KIAA1048), mRNA
NM 014911	Homo sapiens KIAA1042 protein (KIAA1042), mRNA
NM 014947	Homo sapiens KIAA1041 protein (KIAA1041), mRNA
NM 014923	Homo sapiens KIAA0970 protein (KIAA0970), mRNA
NM 015310	Homo sapiens KIAA0942 protein (KIAA0942), mRNA
NM 015057	Homo sapiens KIAA0916 protein (KIAA0916), mRNA
NM 014944	Homo sapiens KIAA0911 protein (KIAA0911), mRNA
NM 014961	Homo sapiens KIAA0871 protein (KIAA0871), mRNA
NM 014941	Homo sapiens KIAA0852 protein (KIAA0852), mRNA
NM 015376	Homo sapiens KIAA0846 protein (KIAA0846), mRNA
NM 014715	Homo sapiens KIAA0712 gene product (KIAA0712), mRNA
NM 014871	Homo sapiens KIAA0710 gene product (KIAA0710), mRNA
NM 014799	Homo sapiens hephaestin (HEPH), mRNA
NM_014678	Homo sapiens KIAA0685 gene product (KIAA0685), mRNA
NM_014011	Homo sapiens KIAA0671 gene product (KIAA0671), mRNA
NM_014741	Homo sapiens KIAA0652 gene product (KIAA0652), mRNA
NM_014662	Homo sapiens KIAA0645 gene product (KIAA0645), mRNA
NM_014838	Homo sapiens KIAA0637 gene product (KIAA0637), mRNA
NM_014774	Homo sapiens KIAA0494 gene product (KIAA0494), mRNA
NM_014870	Homo sapiens KIAA0478 gene product (KIAA0478), mRNA
NM_014856	Homo sapiens KIAA0476 gene product (KIAA0476), mRNA
NM_014864	Homo sapiens KIAA0475 gene product (KIAA0475), mRNA
NM_014857	Homo sapiens KIAA0471 gene product (KIAA0471), mRNA
NM_014812	Homo sapiens KIAA0470 gene product (KIAA0470), mRNA
NM_014826	Homo sapiens KIAA0451 gene product (KIAA0451), mRNA
NM_014675	Homo sapiens KIAA0445 gene product (KIAA0445), mRNA
NM_014751	Homo sapiens KIAA0429 gene product (KIAA0429), mRNA
NM_014724	Homo sapiens KIAA0426 gene product (KIAA0426), mRNA
NM_014684	Homo sapiens KIAA0373 gene product (KIAA0373), mRNA
NM_014809	Homo sapiens KIAA0319 gene product (KIAA0319), mRNA
NM_014727	Homo sapiens KIAA0304 gene product (KIAA0304), mRNA
NM_014807	Homo sapiens KIAA0285 gene product (KIAA0285), mRNA
NM_014767	Homo sapiens KIAA0275 gene product (KIAA0275), mRNA
NM_014785	Homo sapiens KIAA0258 gene product (KIAA0258), mRNA

	CONTAINED TO THE PROPERTY OF T
NM_015153	Homo sapiens KIAA0244 protein (KIAA0244), mRNA
NM_014747	Homo sapiens KIAA0237 gene product (KIAA0237), mRNA
NM_014873	Homo sapiens KIAA0205 gene product (KIAA0205), mRNA
NM_014846	Homo sapiens KIAA0196 gene product (KIAA0196), mRNA
NM_014738	Homo sapiens KIAA0195 gene product (KIAA0195), mRNA
NM_014640	Homo sapiens KIAA0173 gene product (KIAA0173), mRNA
NM_014666	Homo sapiens KIAA0171 gene product (KIAA0171), mRNA
NM_014641	Homo sapiens KIAA0170 gene product (KIAA0170), mRNA
NM_014737	Homo sapiens Ras association (RalGDS/AF-6) domain family 2 (RASSF2), mRNA
NM 014770	Homo sapiens KIAA0167 gene product (KIAA0167), mRNA
NM 014739	Homo saniens KIA A0164 gene product (KIAA0164), mRNA
NM 014865	Homo sapiens chromosome condensation-related SMC-associated protein 1
14141_014603	(KIAA0159), mRNA
NM_014748	Homo sapiens KIAA0064 gene product (KIAA0064), mRNA
NM 014876	Homo sapiens KIAA0063 gene product (KIAA0063), mRNA
NM 014764	Homo sapiens DAZ associated protein 2 (DAZAP2), mRNA
NM 014704	Homo sapiens KIAA0042 gene product (KIAA0042), mRNA
NM 014642	Homo saniens KIA A0036 gene product (KIAA0036), mRNA
NM 015340	Homo sapiens leucyl-tRNA synthetase, mitochondrial (KIAA0028), mRNA
NM 014634	Homo sapiens KIAA0015 gene product (KIAA0015), mRNA
NM 014783	Homo sapiens KIAA0013 gene product (KIAA0013), mRNA
NM 014008	Homo saniens IM1 protein (IM1), mRNA
NM 014066	Homo sapiens HT002 protein; hypertension-related calcium-regulated gene
11111_014000	(HT002), mRNA
NM 014154	Homo sapiens HSPC056 protein (HSPC056), mRNA
NM 014153	Homo sapiens HSPC055 protein (HSPC055), mRNA
NM 014150	Homo sapiens HSPC052 protein (HSPC052), mRNA
NM 014149	Homo sapiens HSPC049 protein (HSPC049), mRNA
NM 014029	Homo sapiens HSPC022 protein (HSPC022), mRNA
NM 014027	Homo sapiens HSPC018 protein (HSPC018), mRNA
NM 014019	Homo sapiens HSPC009 protein (HSPC009), mRNA
NM 015372	Homo sapiens hypothetical protein (HSN44A4A), mRNA
NM 015343	Homo saniens hypothetical protein (HSA011916), mRNA
NM_014063	Homo sapiens src homology 3 domain-containing protein HIP-55 (HIP-55),
NM_014003	mRNA
NM 014052	Homo sapiens GW128 protein (GW128), mRNA
NM 014888	Homo sapiens predicted osteoblast protein (GS3786), mRNA
NM 014030	Homo sapiens G protein-coupled receptor kinase-interactor 1 (GIT1), mRNA
NM 014077	Homo sapiens DKFZP586O0120 protein (DKFZP586O0120), mRNA
NM 015425	Homo sapiens DKFZP586M0122 protein (DKFZP586M0122), mRNA
NM 015456	Homo sapiens DKFZP586B0519 protein (DKFZP586B0519), mRNA
NM 015393	Homo sapiens DKFZP564O0823 protein (DKFZP564O0823), mRNA
NM 015421	Homo sapiens DKFZP564K2062 protein (DKFZP564K2062), mRNA
NM 015421	Homo sapiens DKFZP564B167 protein (DKFZP564B167), mRNA
NM 015527	Homo sapiens DKFZP434P1750 protein (DKFZP434P1750), mRNA
NM_015458	Homo sapiens DKFZP434K171 protein (DKFZP434K171), mRNA
NM_015438	Homo sapiens N-acetylglucosamine-phosphate mutase (AGM1), mRNA
NM_015434 NM_015699	T150 4 10 2) DNIA
NM 015697	
NM_015702	Itomo suprems mypomonous protess. (

27.6.01.5502	COLOG WAY (COLOG) DNA
NM_015703	Homo sapiens CGI-96 protein (CGI-96), mRNA
NM_015380	Homo sapiens CGI-51 protein (CGI-51), mRNA
NM_014143	Homo sapiens B7-H1 protein (B7-H1), mRNA
NM_014062	Homo sapiens ART-4 protein (ART-4), mRNA
NM_014596	Homo sapiens zinc ribbon domain containing, 1 (ZNRD1), mRNA
NM_014519	Homo sapiens zinc finger protein 232 (ZNF232), mRNA
NM_014437	Homo sapiens zinc/iron regulated transporter-like (ZIRTL), mRNA
NM_015363	Homo sapiens zinc finger, imprinted 2 (ZIM2), mRNA
NM_014232	Homo sapiens vesicle-associated membrane protein 2 (synaptobrevin 2) (VAMP2), mRNA
NM_014233	Homo sapiens upstream binding transcription factor, RNA polymerase I (UBTF), mRNA
NM 014235	Homo sapiens ubiquitin-like 4 (UBL4), mRNA
NM 014383	Homo sapiens testis zinc finger protein (TZFP), mRNA
NM_014547	Homo sapiens tropomodulin 3 (ubiquitous) (TMOD3), mRNA
NM 014548	Homo sapiens tropomodulin 2 (neuronal) (TMOD2), mRNA
NM 014464	Homo sapiens tubulointerstitial nephritis antigen (TIN-AG), mRNA
NM 014258	Homo sapiens synaptonemal complex protein 2 (SYCP2), mRNA
NM 014370	Homo sapiens serine/threonine kinase 23 (STK23), mRNA
NM 014264	Homo sapiens serine/threonine kinase 18 (STK18), mRNA
NM 014467	Homo sapiens sushi-repeat protein (SRPUL), mRNA
NM 014230	Homo sapiens signal recognition particle 68kD (SRP68), mRNA
NM 014320	Homo sapiens putative heme-binding protein (SOUL), mRNA
NM 014426	Homo sapiens sorting nexin 5 (SNX5), mRNA
NM_014311	Homo sapiens single-strand selective monofunctional uracil DNA glycosylase (SMUG1), mRNA
NM_014270	Homo sapiens solute carrier family 7 (cationic amino acid transporter, y+ system), member 9 (SLC7A9), mRNA
NM_014252	Homo sapiens solute carrier family 25 (mitochondrial carrier; ornithine transporter) member 15 (SLC25A15), nuclear gene encoding mitochondrial protein, mRNA
NM 014251	Homo sapiens solute carrier family 25, member 13 (citrin) (SLC25A13), mRNA
NM 014442	Homo sapiens sialic acid binding Ig-like lectin 8 (SIGLEC8), mRNA
NM 014521	Homo sapiens SH3-domain binding protein 4 (SH3BP4), mRNA
NM 014554	Homo sapiens sentrin/SUMO-specific protease (SENP1), mRNA
NM 014563	Homo sapiens spondyloepiphyseal dysplasia, late (SEDL), mRNA
NM_014191	Homo sapiens sodium channel, voltage gated, type VIII, alpha polypeptide (SCN8A), mRNA
NM_014139	Homo sapiens sodium channel, voltage-gated, type XII, alpha polypeptide (SCN12A), mRNA
NM 014363	Homo sapiens spastic ataxia of Charlevoix-Saguenay (sacsin) (SACS), mRNA
NM_014285	Homo sapiens homolog of Yeast RRP4 (ribosomal RNA processing 4), 3'-5'-exoribonuclease (RRP4), mRNA
NM_014496	Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 6 (RPS6KA6), mRNA
NM 014245	Homo sapiens ring finger protein 7 (RNF7), mRNA
NM 014372	Homo sapiens ring finger protein 11 (RNF11), mRNA
NM 014314	Homo sapiens RNA helicase (RIG-I), mRNA
NM 014470	Homo sapiens GTP-binding protein (RHO6), mRNA
NM 014248	Homo sapiens ring-box 1 (RBX1), mRNA
NM 014226	Homo sapiens renal tumor antigen (RAGE), mRNA
NM 014488	Homo sapiens RAB30, member RAS oncogene family (RAB30), mRNA
TAIAT 014400	Tionio sapiens tendo, member tend oneogene tennis (tendos), metar

VM 014353	Homo sapiens RAB26, member RAS oncogene family (RAB26), mRNA
2 4 4 4 6	rr alustorin like 1 (retinal) (C.L.) L. IIIINA
VM_015725	Homo sapiens chisterni-like i (remany (ed-2)). Homo sapiens photoreceptor outer segment all-trans retinol dehydrogenase
- 1	~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~
NM_005973	(PRRDH), mRNA Homo sapiens papillary renal cell carcinoma (translocation-associated) (PRCC),
	DATA
NM 014337	Homo sapiens peptidylprolyl isomerase (cyclophilin)-like 2 (PPIL2), mRNA
NM_014348	Homo sapiens similar to rat integral membrane glycoprotein POM121
014310	(POM121L1), mRNA
NM 015720	Home conjens endoglycan (PODLX2), mRNA
NM 014386	Homo saniens polycystic kidney disease 2-like 2 (PKD2L2), IIINNA
NM 014390	TI remines EDNA 2 co-activator (100kl)) (D100), IRKINA
NM 014321	Homo sapiens origin recognition complex, subunit 6 (yeast homolog)-like
MM_014321	CONCIA DATA
ND 4 014566	Homo sapiens olfactory receptor, family 1, subfamily D, member 5 (OR1D5),
NM_014566	TONEA
ND 6 014565	Homo sapiens olfactory receptor, family 1, subfamily A, member 1 (OR1A1),
NM_014565	mRNA
27.6.01.4250	Homo sapiens POU transcription factor (OCT11), mRNA
NM_014352	Homo sapiens odorant-binding protein 2B (OBP2B), mRNA
NM_014581	Try assess advent hinding protein 2A (OBP2A), mRNA
NM_014582	Homo sapiens oddrant-bilding protein 217 (222-23)  Homo sapiens nudix (nucleoside diphosphate linked moiety X)-type motif 5
NM_014142	l a momes DNIA
	(NUDT5), mRNA  Homo sapiens nuclear matrix protein NMP200 related to splicing factor PRP19
NM_014502	Homo sapiens nuclear matrix protein fitti 200 totale to 1
	(NMP200), mRNA  Homo sapiens nesca protein (NESCA), mRNA
NM_014328	Homo sapiens NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 8
NM_014222	Homo sapiens NADH denydiogenase (doiquinone) 1 stylla 1 mp. NA
	(19kD, PGIV) (NDUFA8), mRNA
NM_015678	Homo sapiens neurobeachin (NBEA), mRNA
NM_014461	Homo sapiens contactin 6 (CNTN6), mRNA  Homo sapiens contactin 6 (CNTN6), mRNA
NM_014520	Homo sapiens MYB binding protein (P160) la (MYBBP1A), mRNA  Homo sapiens MYB binding protein (P160) la (MYBBP1A), mRNA
NM_014221	Homo sapiens mature T-cell proliferation 1 (MTCP1), mRNA  Homo sapiens mature T-cell proliferation 2 (MEAP3), mRNA
NM_005927	Homo sapiens microfibrillar-associated protein 3 (MFAP3), mRNA
NM_014623	Homo sapiens male-enhanced antigen (MEA), mRNA
NM 014462	Homo sapiens Lsm1 protein (LSM1), mRNA
NM_014622	Homo sapiens loss of heterozygosity, 11, chromosomal region 2, gene A
\	(I OH11CR2A) mRNA
NM 014240	Homo sapiens LIM domains containing 1 (LIMD1), mRNA
NM_014564	Homo sapiens LIM homeobox protein 3 (LHX3), mkNA
NM 014553	Homo saniens I BP protein (LBP-9), mRNA
NM_014387	Trans against linker for activation of T cells (LA1), mKNA
NM 014379	Home conjens neuronal notassium channel alpha subunit (KV8.1), IIKNA
NM_014514	Homo sapiens killer cell immunoglobulin-like receptor, three domains, short
NM_014513	Homo sapiens killer cell immunoglobulin-like receptor, two domains, short
1.1.1_01.010	
NM_014512	Homo sapiens killer cell immunoglobulin-like receptor, two domains, short
14141_014215	extenlermic tail 1 (KIR2DS1) mRNA
NM 014511	Homo sapiens killer cell immunoglobulin-like receptor, two domains, long
14147_014211	outonlasmic tail 3 (KIR2DI 3) mRNA
NM_014219	
I INIVI UIHAID	cytoplasmic tail, 2 (KIR2DL2), mRNA

NM_014218	Homo sapiens killer cell immunoglobulin-like receptor, two domains, long cytoplasmic tail, 1 (KIR2DL1), mRNA
ND4 014765	
NM_014765	Homo sapiens translocase of outer mitochondrial membrane 20 (yeast) homolog (KIAA0016), mRNA
NM 014406	Homo sapiens potassium large conductance calcium-activated channel,
-	subfamily M, beta member 3-like (KCNMB3L), mRNA
NM 014407	Homo sapiens potassium large conductance calcium-activated channel,
-	subfamily M beta member 3 (KCNMB3), mRNA
NM 014216	Homo sapiens inositol 1,3,4-triphosphate 5/6 kinase (ITPK1), mRNA
NM 014425	Homo sapiens inversin (INVS), mRNA
NM 014214	Homo sapiens inositol(myo)-1(or 4)-monophosphatase 2 (IMPA2), mRNA
NM 014271	Homo sapiens interleukin 1 receptor accessory protein-like 1 (IL1RAPL1),
	mRNA
NM_014339	Homo sapiens interleukin 17 receptor (IL17R), mRNA
NM_014443	Homo sapiens interleukin 17B (IL17B), mRNA
NM_014333	Homo sapiens immunoglobulin superfamily, member 4 (IGSF4), mRNA
NM_014262	Homo sapiens hypothetical protein B (HSU47926), mRNA
NM_014424	Homo sapiens heat shock 27kD protein family, member 7 (cardiovascular)
	(HSPB7), mRNA
NM_014473	Homo sapiens putative dimethyladenosine transferase (HSA9761), mRNA
NM 015370	Homo sapiens hypothetical protein (HS747E2A), mRNA
NM 015371	Homo sapiens hypothetical protein (HS322B1A), mRNA
NM 014345	Homo sapiens endocrine regulator (HRIHFB2436), mRNA
NM 014255	Homo sapiens transmembrane protein 4 (TMEM4), mRNA
NM 014257	Homo sapiens CD209 antigen-like (CD209L), mRNA
NM 014213	Homo sapiens homeo box D9 (HOXD9), mRNA
NM 014620	Homo sapiens homeo box C4 (HOXC4), mRNA
NM 014212	Homo sapiens homeo box C11 (HOXC11), mRNA
NM 014260	Homo sapiens HLA class II region expressed gene KE2 (HKE2), mRNA
NM 014356	Homo sapiens HGC6.2 protein (HGC6.2), mRNA
NM 014354	Homo sapiens HGC6.1.1 protein (HGC6.1.1), mRNA
NM 014571	Homo canions heim/onhances of all to all to 1 APPRIL (1977)
	Homo sapiens hairy/enhancer-of-split related with YRPW motif-like (HEYL), mRNA
NM_014606	Homo sapiens hect domain and RLD 3 (HERC3), mRNA
NM_015726	Homo sapiens H326 (H326), mRNA
NM_014619	Homo sapiens glutamate receptor, ionotropic, kainate 4 (GRIK4), mRNA
NM_014626	Homo sapiens G protein-coupled receptor 58 (GPR58), mRNA
NM_014627	Homo sapiens G protein-coupled receptor 57 (GPR57), mRNA
NM_014498	Homo sapiens type II Golgi membrane protein (GPP130), mRNA
NM_014373	Homo sapiens putative G protein-coupled receptor (GPCR150), mRNA
NM_014236	Homo sapiens glyceronephosphate O-acyltransferase (GNPAT), mRNA
NM_015710	Homo sapiens glioma tumor suppressor candidate region gene 2 (GLTSCR2), mRNA
NM_015711	Homo sapiens glioma tumor suppressor candidate region gene 1 (GLTSCR1), mRNA
NM_015715	Homo sapiens group III secreted phospholipase A2 (GIII-SPLA2), mRNA
NM 014291	Homo sapiens glycine C-acetyltransferase (2-amino-3-ketobutyrate coenzyme A
	ligase) (GCAT), mRNA
NM_014364	Homo sapiens glyceraldehyde-3-phosphate dehydrogenase, testis-specific (GAPDS), mRNA
NM_015714	Homo sapiens putative lymphocyte G0/G1 switch gene (G0S2), mRNA
NM 014489	Homo sapiens FGF receptor activating protein 1 (FRAG1), mRNA
1111 014407	romo sapiens for receptor activating protein 1 (FRAG1), mRNA

	G 11 11 (proton coupled divalent metal ion
IM_014585	Homo sapiens solute carrier family 11 (proton-coupled divalent metal ion
	transporters), member 3 (SLC11A3), mRNA  Homo sapiens putative secreted ligand homologous to fjx1 (FJX1), mRNA
VM_014344	Homo sapiens putative secreted figure formily 7 (FII 1 (7FTA)), mRNA
NM_014439	Homo sapiens Interleukin-1 Superfamily z (FIL1(ZETA)), mRNA  Homo sapiens Interleukin-1 Superfamily z (FIL1(ZETA)), mRNA
NM_014440	Homo sapiens Interleukin-1 Superfamily 1 (FIL1(EPSILON)), mRNA  Homo sapiens Interleukin-1 Superfamily 1 (FIL1) mRNA
NM_014438	Homo sapiens Interleukin-1 Superfamily e (FIL1), mRNA
NM_014210	Homo sapiens ecotropic viral integration site 2A (EVI2A), mRNA
NM_014355	Homo sapiens enolase alpha, lung-specific (ENO1B), mRNA
NM_014600	Homo sapiens EH-domain containing 3 (EHD3), mRNA
NM 014601	Homo sapiens EH-domain containing 2 (EHD2), mRNA
NM 014503	Homo sapiens down-regulated in metastasis (DRIM), mRNA
NM 014549	Try DVE7-424D211 protein (1)K FZP434P211), IIINNA
NM 014388	Homo saniens novel putative protein similar to YIL091C yeast hypothetical 64
	1-D protein from SGA1-KTR7 (D.1434014.5), mKNA
NM_014618	Homo sapiens deleted in bladder cancer chromosome region candidate 1
11212_0	(DRCCR1) mRNA
NM 014392	VI and a reason specific protein (D4S234E), mRNA
NM 004389	Homo sapiens catenin (cadherin-associated protein), aipiia 2 (CINIVA2), met vi
NM 014343	Thomas ganians claudin 15 (CLDN15), mRNA
NM 014887	Homo sapiens hypothetical protein from BCRA2 region (CG003), IIICVA
NM 014207	Trans ganiens CD5 antigen (n56-62) (CD5), mKNA
NM 014335	Home geniens chromosome 15 open reading frame 3 (C150RF3), mRNA
NM 014206	Home series chromosome 11 open reading frame 10 (C1101110), IIIXIVA
NM 014453	There are some nutative breast adenocarcinoma marker (32kD) (BC-2), include
NM_014382	Homo sapiens ATPase, Ca++ transporting, type 2C, member 1 (ATP2C1),
14141_014302	DNA
NM_014570	Homo sapiens ADP-ribosylation factor GTPase activating protein 1
14141_014570	(ADEGAPI) mRNA
NM_014278	Homo saniens heat shock protein (hspl 10 family) (APG-1), mRNA
NM 014495	Trans conjent angionojetin-like 3 (AN(iP1L3), MKNA
NM_004037	Homo sapiens adenosine monophosphate deaminase 2 (isoform L) (AMPD2),
14141_004037	mPNA
NM 014324	Homo saniens alpha-methylacyl-CoA racemase (AMACR), mRNA
NM 014476	Homo sapiens alpha-actinin-2-associated LIM protein (ALP), mRNA
	Try are serious ATT1 fixed gene from 5031 (AF5O31), MKNA
NM_014423	Homo sapiens endogenous retroviral family W, env(C7), member 1 (syncytin)
NM_014590	(ERVWE1), mRNA
27.6 014496	The regions neuronal thread protein (AD7C-NTP), mRNA
NM_014486	Homo sapiens acyl-Coenzyme A dehydrogenase family, member 8 (ACAD8),
NM_014384	DATA
3 D. 4. 014074	Home capiens Alu-hinding protein with zinc finger domain (ABP/ZF), filkINA
NM_014274	Home conjens gamma tubulin ring complex protein (70p gene) (70r), indiva
NM_014444	
NM_007082	DNIA
27 6 010410	mRNA Homo sapiens RAB, member of RAS oncogene family-like 2A (RABL2A),
NM_013412	1 ( a resist variant 1 mPNΔ
	transcript variant 1, mRNA  Homo sapiens peroxisome proliferative activated receptor, alpha (PPARA),
NM_005036	Homo sapiens peroxisonic promerative activated reserving a few property of the contractive activated reserving and the contractive activated reserving a few property of the contractive activated reserving and a few property of the contractive activated reserving a
	mRNA Homo sapiens deiodinase, iodothyronine, type II (DIO2), transcript variant 2,
NM_000793	Homo sapiens defodinase, fododifyrolinie, type if (2202), and a party
	mRNA Homo sapiens deiodinase, iodothyronine, type II (DIO2), transcript variant 1,
NM_013989	
1	mRNA

A to,
to,
to,
,
)2),
A1),
H1),
C1),
H3),
H2),
RNA
A
D1)
d
d
d 
d 

VM 000487	Homo sapiens arylsulfatase A (ARSA), mRNA Homo sapiens small nuclear ribonucleoprotein D2 polypeptide (16.5kD)
JM_004597	Homo sapiens small nuclear ribonucleoprotein bz posypopado
_ [	(CNID DID2) mR NA
M 006194	Homo sapiens paired box gene 9 (PAX9), mRNA
	Homo sapiens NME7 (NME7), mRNA
NM_012476	Homo sapiens NME7 (NME7), med at Homo sapiens ventral anterior homeobox 2 (VAX2), mRNA
NM 012253	Homo sapiens transketolase-like 1 (TKTL1), mRNA  Homo sapiens transketolase-like 1 (TKTL1), mRNA
NM 012268	Homo sapiens transketolase-like I (TKTLI), industry Homo sapiens similar to vaccinia virus HindIII K4L ORF (HU-K4), mRNA Homo sapiens similar to vaccinia virus HindIII K4L ORF (HU-K4), mRNA
NM 002017	Home conjens Friend leukemia virus integration i (1 Lii), integratio
NM 006769	
NM_002260	Homo sapiens LIM domain only 4 (LMO4), interview Homo sapiens killer cell lectin-like receptor subfamily C, member 2 (KLRC2),
_	
NM_005317	mRNA  Homo sapiens granzyme M (lymphocyte met-ase 1) (GZMM), mRNA  Homo sapiens granzyme M (lymphocyte met-ase 1) (GZMM), mRNA
NM 004417	
NM 012125	Transport cholinergic recentor, muscarillic 5 (CIRCVIS); MILE
NM 001236	Home seniens carbonyl reductase 3 (CBR3), IIIKNA
NM 013343	TT mains NAG 7 protein (NA(1-/), IIIRNA
NM 013344	Tr
NM 013236	TY
NM 013380	Try gine finger protein 228 (ZNF220), illicity
NM 013362	Try and gine finger protein 225 (ZNF225), Illiniva
NM 013398	II amo geniens zinc finger protein 224 (ZNF224), mititi
NM 013361	Theme conjens zinc finger protein 223 (ZNF223), IIIXVII
NM 013360	Home geniens zinc finger protein 222 (ZNF222), IIIXIVA
NM 013359	Hame conjens gine finger protein 221 (ZNF221), Hikivi
NM 013250	II was remiens gine finger protein 215 (ZNF215), IIINVI
NM 013249	Finger protein //4 (//NFZ/4/, IIII/1/12)
NM 013249	Homo saniens zinc finger protein 180 (HHZ108) (Zivi 100); indeed
NM 013230	Homo sapiens interleukin 19 (IL19), mRNA
NM 013403	TY services ginedin (ZIN) mRNA
	TV are comions are R lymphocyte gene 3 (VFREDS), mic 12
NM_013378	
NM_013270	Homo sapiens testes-specific protease 30 (18130), interest.  Homo sapiens thyrotropin-releasing hormone degrading ectoenzyme (TRHDE),
NM_013381	
NR 6 012215	mRNA Homo sapiens transmembrane phosphatase with tensin homology (TPTE),
NM_013315	
NM 013353	The serious tropomodulin 4 (muscle) (TMOD4), mRNA
NM_013390	Homo saniens transitional enthelia response protein (TERET); Manuel
NM_013319	
NM_013254	
NM_013309	DATA
27.6 012256	· · · · · · · · · · · · · · · · · · ·
NM 013356	TT
NM_013257	Homo saniens CDK4-binding protein p34SEII (SEII), IIICUL
NM_013370	
NM_013243	
NM_01335	
	DADA interacting protein (rabin 1)-like I (NADJILI), miles
NM_01340	
NM_01323	- liferetize activated tecephol. gailling, oback table
NM_01326	(PPARGC1), mRNA

NM_013268	Homo sapiens placental protein 13 (PP13), mRNA
NM 013382	Homo sapiens putative protein O-mannosyltransferase (POMT2), mRNA
NM 013232	Homo sapiens programmed cell death 6 (PDCD6), mRNA
NM 013397	Homo sapiens over-expressed breast tumor protein (OBTP), mRNA
NM_013389	Homo sapiens NPC1 (Niemann-Pick disease, type C1, gene)-like 1 (NPC1L1), mRNA
NM 013326	Homo sapiens colon cancer-associated protein Mic1 (MIC1), mRNA
NM 013238	Homo sapiens DNAJ domain-containing (MCJ), mRNA
NM 013269	Homo sapiens lectin-like NK cell receptor (LLT1), mRNA
NM_013289	Homo sapiens killer cell immunoglobulin-like receptor, three domains, long cytoplasmic tail, 1 (KIR3DL1), mRNA
NB4 012211	Homo sapiens insulin upstream factor 1 (IUF1), mRNA
NM_013311	Homo sapiens interleukin 17C (IL17C), mRNA
NM_013278	Homo sapiens (clone PWHLC2-24) myosin light chain 2 (HUMMLC2B),
NM_013292	mRNA
NM_013288	Homo sapiens DNA binding protein for surfactant protein B (HUMBINDC), mRNA
NM 013244	Homo sapiens UDP-N-acetylglucosamine:a-1,3-D-mannoside beta-1,4-N-
	acetylglucosaminyltransferase IV-homolog (HGNT-IV-H), mRNA
NM 013264	Homo sapiens gonadotropin-regulated testicular RNA helicase (GRTH), mRNA
NM_013281	Homo sapiens fibronectin leucine rich transmembrane protein 3 (FLRT3),
	mRNA (FI PT2)
NM_013231	Homo sapiens fibronectin leucine rich transmembrane protein 2 (FLRT2), mRNA
NM_013241	Homo sapiens FH1/FH2 domain-containing protein (FHOS), mRNA
NM_013342	Homo sapiens TCF3 (E2A) fusion partner (in childhood Leukemia) (TFPT), mRNA
NM_013246	Homo sapiens cardiotrophin-like cytokine; neurotrophin-1/B-cell stimulating factor-3 (CLC), mRNA
NM_013372	Homo sapiens cysteine knot superfamily 1, BMP antagonist 1 (CKTSF1B1), mRNA
NM_013327	Homo sapiens CGI-56 protein (CGI-56), mRNA
NM_013230	Homo sapiens CD24 antigen (small cell lung carcinoma cluster 4 antigen) (CD24), mRNA
NM 013276	Homo sapiens carbohydrate kinase-like (CARKL), mRNA
NM 013399	Homò sapiens chromosome 16 open reading frame 5 (C16orf5), mRNA
NM 006765	Homo sapiens Putative prostate cancer tumor suppressor (N33), mRNA
NM 006792	Homo sapiens mortality factor 4 (MORF4), mRNA
NM_000397	Homo sapiens cytochrome b-245, beta polypeptide (chronic granulomatous disease) (CYBB), mRNA
NM 005098	Homo sapiens musculin (activated B-cell factor-1) (MSC), mRNA
NM 006144	Homo sapiens granzyme A (granzyme 1, cytotoxic T-lymphocyte-associated
	serine esterase 3) (GZMA), mRNA
NM 002047	Homo sapiens glycyl-tRNA synthetase (GARS), mRNA
NM 004405	Homo sapiens distal-less homeo box 2 (DLX2), mRNA
NM 004371	Homo sapiens coatomer protein complex, subunit alpha (COPA), mRNA
NM 005181	Homo sapiens carbonic anhydrase III, muscle specific (CA3), mRNA
NM 001663	Homo sapiens ADP-ribosylation factor 6 (ARF6), mRNA
NM 001662	Homo sapiens ADP-ribosylation factor 5 (ARF5), mRNA
NM 001660	Homo sapiens ADP-ribosylation factor 4 (ARF4), mRNA
NM 001658	Homo sapiens ADP-ribosylation factor 1 (ARF1), mRNA
NM_000492	Homo sapiens cystic fibrosis transmembrane conductance regulator, ATP-
11112 000 772	1 22000 25 27 27 27 27 27 27 27 27 27 27 27 27 27

	binding cassette (sub-family C, member 7) (CFTR), mRNA
VM 003560	Homo sapiens phospholipase A2, group VI (cytosolic, calcium-independent)
111_003500	
JM 004004	Homo sapiens gap junction protein, beta 2, 26kD (connexin 26) (GJB2), IIIKNA
VM 005198	Homo saniens choline kinase-like (CHKL), Ilikiya
VM 012482	Homo saniens zinc finger protein 281 (ZNF281), mRNA
NM 012256	II-ma conjone gine finger protein 212 (ZNF212), mRNA
NM 012479	Homo seniens tyrosine 3-monooxygenase/tryptopnan 5-monooxygenase
[VIVI_012175	activation protein, gamma polypeptide (YWHAG), mknA
NM 012255	Homo saniens 5'-3' exoribonuclease 2 (XRN2), mRNA
NM 012474	Homo saniens uridine monophosphate kinase (UMPK), mkNA
NM 012473	Homo sapiens thioredoxin, mitochondrial (TXN2), mkNA
NM 012466	Homo sapiens tetraspanin TM4-B (TM4-B), mRNA
NM 012465	Homo sapiens tolloid-like 2 (TLL2), mRNA
NM 012464	Homo saniens tolloid-like 1 (TLL1), mRNA
NM 012290	Homo saniens tousled-like kinase I (ILKI), mKNA
NM 012455	TT CECT homolog (TIC) mRNA
NM 012454	Home conjens T cell lymphoma invasion and metastasis 2 (Traviz), fire vi
NM 012251	Homo saniens transcription factor A, mitochondrial (TFAM), mixta
NM 012451	Try and a symposium of CVN(iR4) mKNA
NM_012448	Homo sapiens synaplogyim 4 (STROKT), meets Homo sapiens signal transducer and activator of transcription 5B (STAT5B),
NWI_012446	mRNA
NM 012447	Hame genions stromal antigen 3 (STAG3), mRNA
NM 012445	Home conjens spondin? extracellular matrix protein (SPONZ), incivit
NM 012443	II are regions gnorm associated antigen 6 (SPAGO), MKNA
NM 012443	Homo sapiens solute carrier family 7 (cationic amino acid transporter, y+
NIVI_U12244	$\frac{1}{1}$ $\frac{1}$
NM_012243	Homo saniens solute carrier family 35 (UDP-N-acetylgiucosamme (UD)
14141_012245	Cu ATA - A
NM_012434	Homo sapiens solute carrier family 17 (anion/sugar transporter), member 5
14141_012451	(SI C17A5) mRNA
NM_012432	Homo sapiens SET domain, bifurcated 1 (SETDB1), mRNA
NM 012427	Homo soniens kallikrein 5 (KLK5), mRNA
NM 012236	TT services core comb on midleg homolog I (SCMH1), IRKNA
NM 012424	Homo sapiens ribosomal protein S6 kinase, 52kD, polypeptide 1 (RPS6KC1),
[NIVI_012421	DNIA
NM_012421	Have coming rearranged I -myc fusion sequence (RLF), mRNA
NM 012415	TY DADA C caregidae homolog ol. D (NAD) 10 11 11 11 12 12 1
NM 012410	Homo sapiens kAD34, S. cerevisiae, nomerce et al. Homo sapiens type I transmembrane receptor (seizure-related protein) (PSK-1),
1411_012-110	mDNA
NM 012409	Homo sapiens prion gene complex, downstream (PRND), mRNA
NM 012402	There's conjugate partner of RAC1 (artantin 2) (POR1), IIIRNA
NM 012400	Tr ranions mhospholinase A2 oroup III) (PLA2G2D), MIKNA
NM 012399	Homo saniens phosphotidylinositol transfer protein, beta (FIII ND), mid 17
NM_012088	Home seniers 6-phosphogluconolactonase (PGLS), mRNA
NM 012395	TI DETAIDE protein kingse (PFIKI), MKNA
NM 012393	
IAIAI_017231	mDNA
NM_012385	Homo seniens no protein (candidate of metastasis 1) (P8), mRNA
	Try
NM_012383 NM_012375	Homo sapiens offictory receptor, family 52, subfamily A, member 1 (OR52A1)
ININI 0172/2	Tromo subjeme orimotory

	10 I C I C I C I C I C I C I C I C I C I
NM_012368	Homo sapiens olfactory receptor, family 2, subfamily C, member 1 (OR2C1), mRNA
NM_012360	Homo sapiens olfactory receptor, family 1, subfamily F, member 8 (OR1F8), mRNA
NM_012352	Homo sapiens olfactory receptor, family 1, subfamily A, member 2 (OR1A2), mRNA
NM_012351	Homo sapiens olfactory receptor, family 10, subfamily J, member 1 (OR10J1), mRNA
NM_012345	Homo sapiens nuclear fragile X mental retardation protein interacting protein 1 (NUFIP1), mRNA
NM_012344	Homo sapiens neurotensin receptor 2 (NTSR2), mRNA
NM_012343	Homo sapiens nicotinamide nucleotide transhydrogenase (NNT), mRNA
NM_012342	Homo sapiens putative transmembrane protein (NMA), mRNA
NM_012337	Homo sapiens nasopharyngeal epithelium specific protein 1 (NESG1), mRNA
NM_012330	Homo sapiens histone acetyltransferase (MORF), mRNA
NM 012064	Homo sapiens major intrinsic protein of lens fiber (MIP), mRNA
NM 012214	Homo sapiens mannosyl (alpha-1,3-)-glycoprotein beta-1,4-N-
_	acetylglucosaminyltransferase, isoenzyme A (MGAT4A), mRNA
NM 012213	Homo sapiens malonyl-CoA decarboxylase (MLYCD), mRNA
NM 012325	Homo sapiens microtubule-associated protein, RP/EB family, member 1
_	(MAPRE1), mRNA
NM 012318	Homo sapiens leucine zipper-EF-hand containing transmembrane protein 1
_	(LETM1), mRNA
NM 012317	Homo sapiens leucine zipper, down-regulated in cancer 1 (LDOC1), mRNA
NM_012314	Homo sapiens killer cell immunoglobulin-like receptor, two domains, short
	cytoplasmic tail, 4 (KIR2DS4), mRNA
NM_012313	Homo sapiens killer cell immunoglobulin-like receptor, two domains, short cytoplasmic tail, 3 (KIR2DS3), mRNA
NM_012312	Homo sapiens killer cell immunoglobulin-like receptor, two domains, short cytoplasmic tail, 2 (KIR2DS2), mRNA
NM_012307	Homo sapiens differentially expressed in adenocarcinoma of the lung (KIAA0987), mRNA
NM 012306	Homo sapiens lifeguard (KIAA0950), mRNA
NM 012302	Homo sapiens latrophilin (KIAA0786), mRNA
NM 012295	Homo sapiens calcineurin binding protein 1 (KIAA0330), mRNA
NM 012288	Homo sapiens TRAM-like protein (KIAA0057), mRNA
NM 012286	Homo sapiens MORF-related gene X (KIAA0026), mRNA
NM_012283	Homo sapiens potassium voltage-gated channel, subfamily G, member 2 (KCNG2), mRNA
NM_012282	Homo sapiens potassium voltage-gated channel, Isk-related family, member 1-like (KCNE1L), mRNA
NM 012278	Homo sapiens integrin beta 1 binding protein (melusin) 2 (ITGB1BP2), mRNA
NM 012211	Homo sapiens integrin, alpha 11 (ITGA11), mRNA
NM 012277	Homo sapiens pancreatic beta cell growth factor (INGAP), mRNA
NM 012275	Homo sapiens interleukin-1 receptor antagonist homolog 1 (IL1HY1), mRNA
NM_012259	Homo sapiens hairy/enhancer-of-split related with YRPW motif 2 (HEY2), mRNA
NM_012258	Homo sapiens hairy/enhancer-of-split related with YRPW motif 1 (HEY1), mRNA
NM 012257	Homo sapiens HMG-box containing protein 1 (HBP1), mRNA
NM_012087	Homo sapiens general transcription factor IIIC, polypeptide 5 (63kD) (GTF3C5), mRNA

NM_012203	Homo sapiens glyoxylate reductase/hydroxypyruvate reductase (GRHPR), mRNA
VM_012202	Homo sapiens guanine nucleotide binding protein (G protein), gamma 3
NM 012084	Homo saniens Glutamate dehydrogenase-2 (GLUD2), mRNA
NM 012191	Homo sapiens putative tumor suppressor (FUS2), mkNA
NM 012185	Homo sapiens forkhead box E2 (FOXE2), mRNA
NM 012183	Homo sapiens forkhead box D3 (FOXD3), mRNA
NM 012153	II and serious Etc homologous factor (EHF), mRNA
NM_012080	Homo sapiens DNA segment, numerous copies, expressed probes (GS1 gene)
	(DXF68S1E), mRNA Homo sapiens double homeobox, 3 (DUX3), mRNA
NM_012148	TY double homeobox 2 (DUX2), mRNA
NM_012147	TI describe describe midulate kinase (thymidylate kinase) (DI YIVIN), IIININA
NM_012145	Homo sapiens deoxytiyilidylate killase (kiyilidylate killase (kiyi
NM_012144	Homo sapiens solute carrier family 25 (mitochondrial carrier; dicarboxylate
NM_012140	transporter), member 10 (SLC25A10), mRNA
	Homo sapiens dimethylarginine dimethylaminohydrolase 1 (DDAH1), mRNA
NM_012137	Homo sapiens leiomodin 1 (smooth muscle) (LMOD1), mRNA
NM_012134	Homo sapiens coatomer protein complex, subunit gamma 2 (COPG2), mRNA
NM_012133	Homo sapiens coatomer protein complex, sabanti gamma 2 (0000)
NM_012132	Homo sapiens claudin 8 (CLDN8), mRNA
NM_012131	Homo sapiens claudin 17 (CLDN17), mRNA
NM_012130	Homo sapiens claudin 14 (CLDN14), mRNA
NM_012129	Homo sapiens claudin 12 (CLDN12), mRNA  Homo sapiens Claudin 12 (CLDN12), mRNA
NM_012127	Homo sapiens Cip1-interacting zinc finger protein (CIZ1), mRNA  Homo sapiens Cip1-interacting zinc finger protein (CIZ1), mRNA
NM_012126	Homo sapiens carbohydrate (N-acetylglucosamine 6-O) sulfotransferase 5 (CHST5), mRNA
NM_012075	Homo sapiens Conserved gene telomeric to alpha globin cluster (CGTHBA), mRNA
NM 012122	Transportant agrhavylesterase 3 (hrain) (CES3), mRNA
NM_012116	Homo sapiens Cas-Br-M (murine) ectropic retroviral transforming sequence c (CBLC), mRNA
NR 6 010112	Homo sapiens carbonic anhydrase XIV (CA14), mRNA
NM_012113	Homo sapiens BUP protein (BUP), mRNA
NM_012071	Home senions systein-rich hydrophobic domain 2 (CHIC2), mKNA
NM_012110	Trans senions brain-specific membrane-anchored protein (BSMAF), IIINIA
NM_012109 NM_012107	Homo sapiens bromodomain containing protein 75 kDa human homolog (Bi 73)
ND 4 010104	mRNA  Homo sapiens beta-site APP-cleaving enzyme (BACE), mRNA
NM_012104	Homo sapiens beta-site APP-cleaving enzyme 2 (BACE2), mRNA
NM_012105	Trame ganiens ancient ubiquitous protein 1 (AUPI), mKNA
NM_012103	The remaining argining glutamic acid dinentide (RE) repeats (RERE), mRNA
NM_012102 NM_012099	Homo sapiens CD3-epsilon-associated protein; antisense to ERCC-1 (ASE-1),
	mRNA (ANGETI 2) mRNA
NM_012098	Homo sapiens angiopoietin-like 2 (ANGPTL2), mRNA
NM_012067	Homo sapiens aldo-keto reductase family 7, member A3 (aflatoxin aldehyde reductase) (AKR7A3), mRNA
NM 012093	Homo sapiens adenylate kinase 5 (AK5), mRNA
NM 012066	Home comions hymothetical protein (20D7-FC4), mKNA
NM 006276	Homo saniens splicing factor, arginine/serine-rich / (35kD) (5FRS/), mRNA
NM 007054	Homo saniens kinesin family member 3A (KIF3A), mRNA
NM 002201	Homo sapiens interferon stimulated gene (20kD) (ISG20), mRNA

Homo sapiens cytosolic acyl coenzyme A thioester hydrolase (HBACH), mRNA
Homo sapiens solute carrier family 9 (sodium/hydrogen exchanger), isoform 3
(SLC9A3), mRNA
Homo sapiens low density lipoprotein-related protein 2 (LRP2), mRNA
Homo sapiens squalene epoxidase (SQLE), mRNA
Homo sapiens plakophilin 4 (PKP4), mRNA
Homo sapiens amine oxidase, copper containing 3 (vascular adhesion protein 1)
(AOC3), mRNA
Homo sapiens tubby like protein 1 (TULP1), mRNA
Homo sapiens mitogen-activated protein kinase 4 (MAPK4), mRNA
Homo sapiens golgi autoantigen, golgin subfamily a, 4 (GOLGA4), mRNA
Homo sapiens brefeldin A-inhibited guanine nucleotide-exchange protein 1
(BIG1), mRNA
Homo sapiens BCL2-associated athanogene 2 (BAG2), mRNA
Homo sapiens anaplastic lymphoma kinase (Ki-1) (ALK), mRNA
Homo sapiens v-akt murine thymoma viral oncogene homolog 2 (AKT2),
mRNA O (1 CTP 2) PNA
Homo sapiens angiotensin receptor 2 (AGTR2), mRNA
Homo sapiens tissue factor pathway inhibitor (lipoprotein-associated coagulation
inhibitor) (TFPI), mRNA
Homo sapiens protein phosphatase 3 (formerly 2B), catalytic subunit, alpha
isoform (calcineurin A alpha) (PPP3CA), mRNA Homo sapiens amelogenin (X chromosome, amelogenesis imperfecta 1)
(AMELX), mRNA
Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 6
(ABCC6), mRNA
Homo sapiens multimerin (MMRN), mRNA
Homo sapiens heat shock 90kD protein 1, beta (HSPCB), mRNA
Homo sapiens putative GR6 protein (GR6), mRNA
Homo sapiens guanine nucleotide binding protein (G protein) alpha 12 (GNA12),
mRNA
Homo sapiens phospholipase A2 receptor 1, 180kD (PLA2R1), mRNA
Homo sapiens pleckstrin homology-like domain, family A, member 1
(PHLDA1), mRNA
Homo sapiens integral type I protein (P24B), mRNA
Homo sapiens nucleoporin-like protein 1 (NLP_1), mRNA
Homo sapiens nidogen 2 (NID2), mRNA
Homo sapiens SH3 domain binding glutamic acid-rich protein (SH3BGR),
mRNA
Homo sapiens replication factor C (activator 1) 5 (36.5kD) (RFC5), mRNA
Homo sapiens activating transcription factor 6 (ATF6), mRNA
Homo sapiens Rho-associated, coiled-coil containing protein kinase 2 (ROCK2),
mRNA
Homo sapiens LIM domain only 2 (rhombotin-like 1) (LMO2), mRNA
Homo sapiens deleted in liver cancer 1 (DLC1), mRNA
Homo sapiens BarH-like homeobox 2 (BARX2), mRNA
Homo sapiens synaptogyrin 3 (SYNGR3), mRNA
Homo sapiens etoposide-induced mRNA (PIG8), mRNA
Homo sapiens natural killer-tumor recognition sequence (NKTR), mRNA  Homo sapiens 5,10-methylenetetrahydrofolate reductase (NADPH) (MTHFR),
mRNA
Homo sapiens potassium intermediate/small conductance calcium-activated

PCT/US03/05028 WO 03/074654

	channel, subfamily N, member 1 (KCNN1), mRNA
	Homo sapiens interphotoreceptor matrix proteoglycan 1 (IMPG1), mRNA
VM 001563	Homo sapiens interphotoreceptor matrix proteogrycan i (Mil 97); Homo sapiens gap junction protein, alpha 5, 40kD (connexin 40) (GJA5), mRNA
VM 005266	Homo sapiens gap junction protein, aipha 3, votes
NM 001874	Homo sapiens carboxypeptidase M (CPM), mRNA Homo sapiens ankyrin-like with transmembrane domains 1 (ANKTM1), mRNA Homo sapiens ankyrin-like with transmembrane domains 1 (TSTA3), mRNA
NM 007332	Homo sapiens ankyrin-like with transmethorane domains (CSTA3), mRNA
000010	Theme conjugate specific transplantation antigent 1352 (1997)
NM 001607	Homo sapiens GDP dissociation innition 2 (GDB), find the Homo sapiens acetyl-Coenzyme A acyltransferase 1 (peroxisomal 3-oxoacyl-Homo acyltransferase 1 (peroxisomal 3-oxoacyl-Homo acyltransferase 1 (peroxisomal 3-oxoacyl-Homo acyltransferase 1 (p
_	Homo sapiens acetyl-Coenzyme A acytransicrase i (perometers). Coenzyme A thiolase) (ACAA1), nuclear gene encoding mitochondrial protein,
NM_003145	mRNA Homo sapiens signal sequence receptor, beta (translocon-associated protein beta)
	(GODO) DATA
NM 000852	Homo sapiens glutathione S-transferase pi (GSTP1), mRNA
NM 000827	
NM_005252	Homo sapiens glutamate receptor, followopie, Fixed Taylor (FOS), Homo sapiens v-fos FBJ murine osteosarcoma viral oncogene homolog (FOS),
141/1_00025	mPNA
NM 005803	Homo sapiens flotillin 1 (FLOT1), mRNA
NM 004459	Home caniens fetal Alzheimer antigen (FALZ), IIIXVA
NM 004081	Homo saniens deleted in azoospermia (DAZ), hikiya
NM 004055	Homo saniens calpain 5 (CAPN5), mRNA
NM 004042	Homo saniens arylsulfatase F (ARSF), mRNA
NM 003085	1 :: 1 = 40 /CN/ B) mkinA
NM 000612	1. 1:1
NM 006995	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
	Homo sapiens butyrophilin, sublating 2, memor 12 (Calcium and DAG-regulated)  Homo sapiens RAS guanyl releasing protein 1 (calcium and DAG-regulated)
NM_005739	(DACCEDE) mPNA
NR 4 006267	Home caniens RAN binding protein 2 (RANBP2), HIRNA
NM_006267	TI comicing DANI binding protein 1 (RANDE 1), IIIXXX
NM_002882	
NM_003884	Homo sapiens p300/CBP-associated factor (FCFF),  Homo sapiens GTP cyclohydrolase I feedback regulatory protein (GCHFR),
NM_005258	7374
27 6 001120	Home senions amino-terminal enhancer of split (AES), mRNA
NM_001130	TI was assisted and supplied the prostate (ACII), illiciti
NM_001099	Homo sapiens actu phosphatase, protein thioesterase 2 (PPT2), mRNA  Homo sapiens palmitoyl-protein thioesterase 2 (PPT2), mRNA
NM_005155	Tyr homes how D3 (H()X1)3), mKNA
NM_006898	Term containing monogyverilase 3 (11403), into 12
NM_006894	Homo sapiens flavin containing monocy general (FEN1), mRNA  Homo sapiens flap structure-specific endonuclease 1 (FEN1), mRNA
NM_004111	
NM_001828	Homo sapiens Charot-Leyden crystal protein (ODO), marginal transducer and activator of transcription 1, 91kD (STAT1),
NM_007315	
	mRNA Homo sapiens NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 9 (22kD
NM_005005	Homo sapiens NADH denydrogenase (dorquinos)
	B22) (NDUFB9), mRNA
NM_003362	
NM_005221	Homo sapiens distal-less homeo box 5 (DLX5), mRNA  Homo sapiens distal-less homeo box 5 (DLX5), mRNA
NM_000479	Homo sapiens anti-Mullerian hormone (AMH), mRNA  Homo sapiens anti-Mullerian hormone (AMH), mRNA
NM 005160	TT TT I I I I I I I I I I I I I I I I I
NM 001619	Try and advanced heta recentor killase I (ADRDICE), indices
NM 001611	Uomo saniens acid phosphatase 5, tartrate resistant (ACI 5), mid 12
NM_003403	Homo saviens YY1 transcription factor (YY1), midva
NM_003793	
NM_001922	Homo sapiens dopachrome tautomerase (dopachionie deta-isomerase, s)
1 , , , , , , , , , , , , , , , , , , ,	related protein 2) (DCT), mRNA

37.6.006410	Homo sapiens 1-acylglycerol-3-phosphate O-acyltransferase 2 (lysophosphatidic
NM_006412	acid acyltransferase, beta) (AGPAT2), mRNA
27.6.000010	Homo sapiens gamma-aminobutyric acid (GABA) A receptor, alpha 5
NM_000810	(GABRA5), mRNA
27.6.000420	Homo sapiens platelet-activating factor acetylhydrolase, isoform Ib, alpha
NM_000430	subunit (45kD) (PAFAH1B1), mRNA
)D ( 002006	Subunit (45KD) (FAFARIDI), mixtx
NM_003006	Homo sapiens selectin P ligand (SELPLG), mRNA
NM_002634	Homo sapiens prohibitin (PHB), mRNA
NM_002410	Homo sapiens mannosyl (alpha-1,6-)-glycoprotein beta-1,6-N-acetyl-glucosaminyltransferase (MGAT5), mRNA
NR 6 000 100	Homo sapiens mannosyl (beta-1,4-)-glycoprotein beta-1,4-N-
NM_002409	acetylglucosaminyltransferase (MGAT3), mRNA
NB ( 002400	Homo sapiens mannosyl (alpha-1,6-)-glycoprotein beta-1,2-N-
NM_002408	acetylglucosaminyltransferase (MGAT2), mRNA
ND ( 000406	Homo sapiens mannosyl (alpha-1,3-)-glycoprotein beta-1,2-N-
NM_002406	acetylglucosaminyltransferase (MGAT1), mRNA
ND 4 000000	Homo sapiens mitogen-activated protein kinase kinase kinase 5 (MAP3K5),
NM_005923	mRNA
ND 6 000005	Homo sapiens isovaleryl Coenzyme A dehydrogenase (IVD), nuclear gene
NM_002225	encoding mitochondrial protein, mRNA
277.6.001.400	Homo sapiens galanin receptor 1 (GALR1), mRNA
NM_001480	Homo sapiens garanti receptor I (GALRI), integral Homo sapiens coagulation factor II (thrombin) receptor (F2R), mRNA
NM_001992	Homo sapiens adenosine A3 receptor (ADORA3), mRNA
NM_000677	Homo sapiens adenosine A3 receptor (ADORAS), mixtor
NM_002969	Homo sapiens mitogen-activated protein kinase 12 (MAPK12), mRNA
NM_001526	Homo sapiens hypocretin (orexin) receptor 2 (HCRTR2), mRNA
NM_003605	Homo sapiens O-linked N-acetylglucosamine (GlcNAc) transferase (UDP-N-
	acetylglucosamine:polypeptide-N-acetylglucosaminyl transferase) (OGT),
	mRNA CD40D alaba 4 subunit of VI A 4
NM_000885	Homo sapiens integrin, alpha 4 (antigen CD49D, alpha 4 subunit of VLA-4
	receptor) (ITGA4), mRNA
NM_003197	Homo sapiens transcription elongation factor B (SIII), polypeptide 1-like
- T C 00 (100	(TCEB1L), mRNA
NM_006183	Homo sapiens neurotensin (NTS), mRNA
NM_002524	Homo sapiens neuroblastoma RAS viral (v-ras) oncogene homolog (NRAS),
	mRNA
NM_002478	Homo sapiens myogenic factor 3 (MYOD1), mRNA
NM_002451	Homo sapiens methylthioadenosine phosphorylase (MTAP), mRNA
NM_002436	Homo sapiens membrane protein, palmitoylated 1 (55kD) (MPP1), mRNA
NM_002377	Homo sapiens MAS1 oncogene (MAS1), mRNA
NM_002305	Homo sapiens lectin, galactoside-binding, soluble, 1 (galectin 1) (LGALS1),
	mRNA (150) 11 (11)
NM_000887	Homo sapiens integrin, alpha X (antigen CD11C (p150), alpha polypeptide)
	(ITGAX), mRNA
NM_000419	Homo sapiens integrin, alpha 2b (platelet glycoprotein IIb of IIb/IIIa complex,
	antigen CD41B) (ITGA2B), mRNA
NM_002203	Homo sapiens integrin, alpha 2 (CD49B, alpha 2 subunit of VLA-2 receptor)
	(ITGA2), mRNA
NM_003637	Homo sapiens integrin, alpha 10 (ITGA10), mRNA
NM_000843	Homo sapiens glutamate receptor, metabotropic 6 (GRM6), mRNA
NM_000838	Homo sapiens glutamate receptor, metabotropic 1 (GRM1), mRNA
NM_000835	Homo sapiens glutamate receptor, ionotropic, N-methyl D-aspartate 2C
_	(GRIN2C), mRNA

PCT/US03/05028 WO 03/074654

<b>W O O O O O O O O O O</b>	1.12 +44 2B
= - 000004	Homo sapiens glutamate receptor, ionotropic, N-methyl D-aspartate 2B
NM_000834	(GRIN2B), mRNA
	(GRIN2B), mRNA Homo sapiens glutamate receptor, ionotropic, N-methyl D-aspartate 2A
NM_000833	(GRIN2A), mRNA
222004	(GRIN2A), mRNA Homo sapiens glutathione peroxidase 3 (plasma) (GPX3), mRNA
NM_002084	Homo sapiens gastrin (GAS), mRNA  Homo sapiens gastrin (GAS), mRNA  atrophy (atrophin-1) (DRPLA),
NM_000805	Homo sapiens gastrin (GAS), mRNA Homo sapiens dentatorubral-pallidoluysian atrophy (atrophin-1) (DRPLA),
NM_001940	Homo sapiens defiator desar p
	mRNA Homo sapiens calumenin (CALU), mRNA Sperm receptor) (ZP3A), mRNA
NM_001219	
NM_007155	
NM_007136	
NM_007250	
NM_007167	
NM_007153	Homo sapiens zinc finger protein 195 (ZNF195), mRNA  Homo sapiens zinc finger protein 195 (ZNF195), mRNA
NM_007152	
NM 007150	Homo sapiens zinc finger protein 105 (ZNF175), mRNA  Homo sapiens zinc finger protein 175 (ZNF175), mRNA
NM 007147	Homo sapiens zinc finger protein 146 (ZNF146), mRNA  Homo sapiens zinc finger protein 146 (ZNF146), mRNA
NM 007145	Homo sapiens zinc finger protein 140 (2/12 14)
NM_007127	Homo sapiens villin 1 (VIL1), mRNA  Homo sapiens ubiquitously transcribed tetratricopeptide repeat gene, Y  AMERICA TRANSCRIPTION OF THE PROPERTY OF THE PROPE
NM 007125	Homo sapiens ubiquitously transcribed tenderto-p
	chromosome (UTY), mRNA
NM_007124	chromosome (UTY), mRNA  Homo sapiens utrophin (homologous to dystrophin) (UTRN), mRNA  Transcription factor 1 (USF1), mRNA
NM_007122	Homo sapiens utrophin (nontologous to eyecr) Homo sapiens upstream transcription factor 1 (USF1), mRNA Homo sapiens UDP glycosyltransferase 2 family, polypeptide B (UGT2B),
NM_007120	Homo sapiens UDP glycosyltransierase 2 lumity, p = 51
14141_00,120	mRNA
NM_007106	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
NM_007118	Homo sapiens triple functional domain (TTPH) mRNA
NM_007117	
NM_007218	
14141_007210	mRNA (TRECTCI) mPNA
NM 007233	
NM 007114	
NM 00711	
NM 00711	
NM_00711	Homo sapiens transcription factor 19 (SC1) (TCF19), mRNA  Homo sapiens transcription factor 19 (SC1) (TCF19), polypeptide 2 (18kD,
NM_00710	Homo saniens transcription elongation factor B (527), F
1VIVI_00710	elongin B) (TCEB2), mRNA elongin B) (TCEB2), mRNA comply 22 (organic cation transporter), member 1-
NM_00710	I I I como caniens solute carrier family 22 (organio data
NM_00/10	like antisense (SLC22A1LS), mRNA
37 6 00716	
NM_00716	
5 5 0071	
NM_00710	Homo sapiens sarcosine dehydrogenase (SARDH), MRNA Homo sapiens splicing factor 3a, subunit 2, 66kD (SF3A2), mRNA Homo sapiens splicing factor 3a, subunit 2, 66kD (SF3A2), mRNA
NM 0071	
NM_0072	Homo sapiens Retina-derived 100 description (REA), mRNA  Homo sapiens B-cell associated protein (REA), mRNA  ONA directed jota (POLI), mRNA
NM_0072	Homo sapiens B-cell associated protein (KE/1), 195 Homo sapiens polymerase (DNA directed) iota (POLI), mRNA Homo sapiens polymerase (DNA directed) iota (POLI), mRNA  195 Homo sapiens B-cell associated protein (NE/1), 195 Homo sapiens polymerase (DNA directed) iota (POLI), mRNA Homo sapiens polymerase (DNA directed) iota (POLI), mRNA Homo sapiens polymerase (DNA directed) iota (POLI), 195 Homo
NM_0071	
NM_0072	Homo sapiens protein tyrosina
	mRNA  Homo sapiens kallikrein 8 (neuropsin/ovasin) (KLK8), mRNA  Homo sapiens kallikrein 8 (neuropsin/ovasin) (KLK8), mRNA
NM_0071	Homo sapiens kallikrein 8 (neuropsin oversin) (120) Homo sapiens protein-O-mannosyltransferase 1 (POMT1), mRNA Homo sapiens protein-O-mannosyltransferase 1 (pompta), mRNA  RNA  RNA  RNA
NM_0071	
NM_0072	Homo sapiens polymerase (DIVA director), g
ı — —	(POLG2), mRNA

	(2)
NM_007254	Homo sapiens polynucleotide kinase 3'-phosphatase (PNKP), mRNA
NM_007221	Homo sapiens polyamine-modulated factor 1 (PMF1), mRNA
NM 007183	Homo sapiens plakophilin 3 (PKP3), mRNA
NM 007169	Homo sapiens phosphatidylethanolamine N-methyltransferase (PEMT), mRNA
NM 007229	Homo sapiens protein kinase C and casein kinase substrate in neurons 2
_	(PACSIN2), mRNA
NM 007190	Homo sapiens Sec23-interacting protein p125 (P125), mRNA
NM 007160	Homo sapiens olfactory receptor, family 2, subfamily H, member 3 (OR2H3),
_	mRNA
NM_007256	Homo sapiens solute carrier family 21 (organic anion transporter), member 9
	(SLC21A9), mRNA
NM 007172	Homo sapiens nucleoporin 50kD (NUP50), mRNA
NM_007103	Homo sapiens NADH dehydrogenase (ubiquinone) flavoprotein 1 (51kD)
	(NDUFV1), mRNA
NM_007181	Homo sapiens mitogen-activated protein kinase kinase kinase kinase 1
	(MAP4K1), mRNA
NM 007230	Homo sapiens mannosidase, alpha, class 1B, member 1 (MAN1B1), mRNA
NM_007164	Homo sapiens mucosal vascular addressin cell adhesion molecule 1
	(MADCAM1), mRNA
NM 007216	Homo sapiens alpha integrin binding protein 63 (KIAA1017), mRNA
NM 007213	Homo sapiens JM4 protein (JM4), mRNA
NM 007102	Homo sapiens guanylate cyclase activator 2B (uroguanylin) (GUCA2B), mRNA
NM 007227	Homo sapiens G protein-coupled receptor 45 (GPR45), mRNA
NM 007275	Homo sapiens lung cancer candidate (FUS1), mRNA
NM 007262	Homo sapiens RNA-binding protein regulatory subunit (DJ-1), mRNA
NM 007166	Homo sapiens Clathrin assembly lymphoid-myeloid leukemia gene (CLTH),
14141_007100	mRNA
NM 007186	Homo sapiens centrosomal protein 2 (CEP2), mRNA
NM 006585	Homo sapiens chaperonin containing TCP1, subunit 8 (theta) (CCT8), mRNA
NM 007185	Homo sapiens trinucleotide repeat containing 4 (TNRC4), mRNA
NM_007220	Homo sapiens carbonic anhydrase VB, mitochondrial (CA5B), nuclear gene
14141_007220	encoding mitochondrial protein, mRNA
NM 007100	Homo sapiens ATP synthase, H+ transporting, mitochondrial F0 complex,
14141_007100	subunit e (ATP5I), mRNA
NM_007231	Homo sapiens solute carrier family 6 (neurotransmitter transporter), member 14
14141_007251	(SLC6A14), mRNA
NM 007203	Homo sapiens A kinase (PRKA) anchor protein 2 (AKAP2), mRNA
NM 007202	Homo sapiens A kinase (PRKA) anchor protein 10 (AKAP10), mRNA
NM 007168	Homo sapiens ATP-binding cassette, sub-family A (ABC1), member 8
NM_00/108	(ABCA8), mRNA
NIM 000506	Homo sapiens coagulation factor II (thrombin) (F2), mRNA
NM_000506	Homo sapiens coagnitation factor if (thromony (12); mkd (1)  Homo sapiens calreticulin (CALR), mRNA
NM_004343	Homo sapiens carrettethin (CALK), inktyA  Homo sapiens heat shock protein, neuronal DNAJ-like 1 (HSJ1), mRNA
NM_006736	Homo sapiens neat snock protein, neuronal DNA3-like 1 (H331), likeva  Homo sapiens erythroid differentiation and denucleation factor 1 (HFL-EDDG1),
NM_006553	
37.6.006004	mRNA
NM_006984	Homo sapiens claudin 10 (CLDN10), mRNA
NM_005502	Homo sapiens ATP-binding cassette, sub-family A (ABC1), member 1
	(ABCA1), mRNA
NM_005809	Homo sapiens peroxiredoxin 2 (PRDX2), mRNA
1 3 75 6 00 60 60	
NM_006977	Homo sapiens zinc finger protein 46 (KUP) (ZNF46), mRNA
NM_006977 NM_006965	Homo sapiens zinc finger protein 46 (KUP) (ZNF46), mRNA Homo sapiens zinc finger protein 24 (KOX 17) (ZNF24), mRNA Homo sapiens zinc finger protein 22 (KOX 15) (ZNF22), mRNA

WO 03/0/4054	
	Homo sapiens zinc finger protein 183 (RING finger, C3HC4 type) (ZNF183),
NM_006978	Homo sapiens zine iniger protein and
	mRNA Homo sapiens uroplakin 3 (UPK3), mRNA
	Homo sapiens uroplakin 1B (UPK1B), Illicara Homo sapiens TATA box binding protein (TBP)-associated factor, RNA
NM_006951	polymerase II, D, 100kD (TAF2D), mRNA
NM_006950	Homo sapiens synapsin I (SYN1), mRNA  Homo sapiens suppressor of white apricot homolog 2 (SWAP2), mRNA  Homo sapiens suppressor of white apricot homolog 2 (STYRP2) mRNA
NM_007056	
NM_006949	Homo sapiens syntaxin binding protein 2 (517622); Homo sapiens stress 70 protein chaperone, microsome-associated, 60kD
NM_006948	Homo sapiens stress /0 protein chaperone, inforcement
- · · -	
NM 006946	(STCH), mRNA  Homo sapiens spectrin, beta, non-erythrocytic 2 (SPTBN2), mRNA  Homo sapiens spectrin, beta, non-erythrocytic 2 (SPRR2B), mRNA
NM 006945	
NM_006944	the second phosphoprojeii 2, 27AD (0112);
NM 007009	
NM 006940	
NM 007017	
NM 006943	
NM_000943	
NM_006942	
NM_006941	Homo sapiens SRY (sex determining region 1) son 15 (Homo sapiens solute carrier family 6 (neurotransmitter transporter, glycine),
NM_006934	
20.5022	member 9 (SLC6A9), mRNA  Homo sapiens solute carrier family 5 (inositol transporters), member 3
NM_006933	(SLC5A3), mRNA
20.6001	(SLC5A3), mRNA Homo sapiens solute carrier family 2 (facilitated glucose transporter), member 3
NM_006931	(SLC2A3), mRNA
NM_006930	Homo sapiens S-phase kinase-associated protein 12 (F)  Homo sapiens splicing factor, arginine/serine-rich 5 (SFRS5), mRNA  Homo sapiens splicing factor, arginine/serine-rich 1 (splicing factor 2, alternate
NM_006925	Homo sapiens splicing factor, arginine/serine-rich 1 (splicing factor 2, alternate Homo sapiens splicing factor, arginine/serine-rich 1 (splicing factor 2, alternate
NM_006924	
	Homo sapiens retinoid X receptor, gamma (RXRG), mRNA  Homo sapiens retinoid X receptor, gamma (RXRG), mRNA  (RYRG), mRNA
NM_006917	Homo sapiens retinoid X receptor, gamma (RARO), MRG 12  Homo sapiens rabphilin 3A-like (without C2 domains) (RPH3AL), mRNA  Homo sapiens rabphilin 3A-like (without C2 domains) (RPH3AL), mRNA  HOMO sapiens rabphilin 3A-like (without C2 domains) (RPH3AL), mRNA
NM_006987	Homo sapiens rabphilin 3A-like (without C2 dollaris) (12 12 13 13 14 15 15 15 15 15 15 15 15 15 15 15 15 15
NM_007055	Homo sapiens polymerase (KIVA) III (BILLE III)
	mRNA (V. linked recessive) (RP2), mRNA
NM_006915	Homo sapiens retinitis pigmentosa 2 (X-Innee Policy Homo sapiens RAR-related orphan receptor B (RORB), mRNA  Homo sapiens RAR-related orphan receptor B (RORB), mRNA
NM_006914	TT
NM 006913	TI and conjent ring finger protein 5 (Kill 5), me
NM 006911	
NM_007043	TYTE I I work binding project ( ) III Day in a second
NM_007033	Homo sapiens HIV-1 rev binding protein 2 (RER1), mRNA  Homo sapiens similar to S. cerevisiae RER1 (RER1), mRNA  Homo sapiens similar to S. cerevisiae RER1 (RER1), mRNA
NM_00708	
14141_00700	mRNA is the 1 glycoprotein 1 (PSG1), mRNA
NM_00690	
NM_00701	
INIVI_00/01	m D N A
3TM 6 00702	DI 6 mrotain (PI 6) mKNA
NM_00702	2 III-ma conjene brain-specific protein p23 alpha (p23);
NM_00703	I Homo copiens myosin IXA (MYO9A), mkiva
NM_00690	IT III and conjens IM5 protein (JM5), miking
NM_00707	Homo sapiens JM27 protein (JM27), mRNA  Homo sapiens JM27 protein (JM27), mRNA
NM 00700	
NM_00689	99   Homo sapiens isocitate denyer g

NM_007031	Homo sapiens heat shock transcription factor 2 binding protein (HSF2BP), mRNA
NM 007011	Homo sapiens putative transmembrane protein (HS1-2), mRNA
NM 006896	Homo sapiens homeo box A7 (HOXA7), mRNA
NM 007045	Homo sapiens FGFR1 oncogene partner (FOP), mRNA
NM 007051	Homo sapiens Fas (TNFRSF6) associated factor 1 (FAF1), mRNA
NM 006979	Homo sapiens HLA class II region expressed gene KE4 (HKE4), mRNA
NM 007015	Homo sapiens chondromodulin I precursor (CHM-I), mRNA
NM 006890	Homo sapiens carcinoembryonic antigen-related cell adhesion molecule 7
NM_000890	(CEACAM7), mRNA
NM_007018	Homo sapiens centrosomal protein 1 (CEP1), mRNA
NM_006889	Homo sapiens CD86 antigen (CD28 antigen ligand 2, B7-2 antigen) (CD86), mRNA
NM 006982	Homo sapiens cartilage paired-class homeoprotein 1 (CART1), mRNA
NM 007058	Homo sapiens calpain 11 (CAPN11), mRNA
NM 006888	Homo sapiens calmodulin 1 (chi 1411), inictia  Homo sapiens calmodulin 1 (phosphorylase kinase, delta) (CALM1), mRNA
	<u> </u>
NM_007047	Homo sapiens butyrophilin, subfamily 3, member A2 (BTN3A2), mRNA  Homo sapiens butyrophilin, subfamily 3, member A1 (BTN3A1), mRNA
NM_007048	
NM_006992	Homo sapiens B7 protein (B7), mRNA
NM_006885	Homo sapiens AT-binding transcription factor 1 (ATBF1), mRNA
NM_007022	Homo sapiens putative tumor suppressor (101F6), mRNA
NM 006697	Homo sapiens cisplatin resistance associated (CRA), mRNA
NM_006826	Homo sapiens tyrosine 3-monooxygenase/tryptophan 5-monooxygenase
NTM 000761	activation protein, theta polypeptide (YWHAQ), mRNA
NM_006761	Homo sapiens tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, epsilon polypeptide (YWHAE), mRNA
NM 006784	Homo sapiens WD repeat domain 3 (WDR3), mRNA
NM 006846	Homo sapiens serine protease inhibitor, Kazal type, 5 (SPINK5), mRNA
NM_006830	Homo sapiens ubiquinol-cytochrome c reductase (6.4kD) subunit (UQCR),
) D ( 00 ( 500	mRNA
NM_006798	Homo sapiens UDP glycosyltransferase 2 family, polypeptide A1 (UGT2A1),
) D 6 006767	mRNA
NM_006757	Homo sapiens troponin T3, skeletal, fast (TNNT3), mRNA
NM_006827	Homo sapiens transmembrane trafficking protein (TMP21), mRNA
NM_006853	Homo sapiens kallikrein 11 (KLK11), mRNA
NM_006811	Homo sapiens tumor differentially expressed 1 (TDE1), mRNA
NM_006756	Homo sapiens transcription elongation factor A (SII), 1 (TCEA1), mRNA
NM_006024	Homo sapiens Tax1 (human T-cell leukemia virus type I) binding protein 1 (TAX1BP1), mRNA
NM 006752	Homo sapiens surfeit 5 (SURF5), mRNA
NM_006819	Homo sapiens stress-induced-phosphoprotein 1 (Hsp70/Hsp90-organizing protein) (STIP1), mRNA
NM 006780	Homo sapiens SMA3 (SMA3), mRNA
NM 006749	Homo sapiens SMAS (SMAS), Inkiva  Homo sapiens solute carrier family 20 (phosphate transporter), member 2
14141_000/49	(SLC20A2), mRNA
NM 006747	Homo sapiens signal-induced proliferation-associated gene 1 (SIPA1), mRNA
NM 006873	Homo sapiens stoned B/TFIIA-alpha/beta-like factor (SALF), mRNA
NM 006788	Homo sapiens ralA binding protein 1 (RALBP1), mRNA
	Homo sapiens receptor-interacting serine-threonine kinase 3 (RIPK3), mRNA
NM_006871	
NM_006867	Homo sapiens RNA-binding protein gene with multiple splicing (RBPMS), mRNA
NM_006743	Homo sapiens RNA binding motif protein 3 (RBM3), mRNA

VM 006868	Homo sapiens RAB31, member RAS oncogene family (RAB31), mRNA
VM_006839	Homo sapiens RAB31, member RAS one ogen and finite filin) (IMMT), Homo sapiens inner membrane protein, mitochondrial (mitofilin) (IMMT),
	DNIA
NM 006812	Homo sapiens amplified in osteosarcoma (OS-9), mRNA
TO COCCEC	Uamo saniens sialidase 3 (membrane sialidase) (NEOS), inicura
NM 006791	Homo saniens MORF-related gene 13 (MRG13), http://
NM 006766	
NM 006804	the series courte remilatory ntoletil telated type 10-75 into 12-2
NM_006770	Homo sapiens macrophage receptor with collagenous structure (Wirksco),
NM_006785	mRNA  Homo sapiens mucosa associated lymphoid tissue lymphoma translocation gene 1 (MALT1), mRNA
NM 006767	like transcriptional regulator, I (LZIKI), many
NM_006840	Homo saniens leukocyte immunoglobulin-like receptor, sublatini b (with 1212
14141_0000	l a week of 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
NM_006866	Homo sapiens leukocyte immunoglobulin-like receptor, sublamily A (with 114)
	Homo sapiens leukocyte immunoglobulin-like receptor, subfamily A (with TM
NM_006863	
27.5.006047	Homo sapiens leukocyte immunoglobulin-like receptor, subfamily B (with TM
NM_006847	1 rows ( 1 - : )
27.5.000005	Homo sapiens leukocyte immunoglobulin-like receptor, subfamily A (without
NM_006865	1 2 /T II D A 3 \ mR N A
ND 4 000064	Homo sapiens leukocyte immunoglobba w BNA
NM_006864	and FFIM domains) member 3 (LILRB3), mRNA
ND 4 006729	TT assigns lymphoid blast crisis oncogene (LDC), nikkyA
NM_006738	Homo sapiens Lysosomal-associated multispanning membrane protein-5
NM_006762	(TADTRAS) DNIA
277 000727	Homo sapiens killer cell immunoglobulin-like receptor, three domains, long
NM_006737	1 . 1 :- 4-31 2 (IZID 2DI 2) mRNA
NTM 006901	Homo sapiens KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein
NM_006801	1 // DEI DI) mRNA
NM 006844	Try (hoctorial acetolactate synthase)-like (IL V DL), illicity
	TT
NM_006858	Homo sapiens interferon-related developmental regulator 2 (IFRD2), mRNA
NM_006764	TI conjent ATP/GTP-hinding protein (HEAB), MRNA
NM_006831	Type
NM_006794	Try and the second symptom protein here had been been been been been been been bee
NM_006783	Homo sapiens gap junction protein, sear o (constant) Homo sapiens FSH primary response (LRPR1, rat) homolog 1 (FSHPRH1),
NM_006733	T DATA
37.4.00(721	Homo sapiens Fukuyama type congenital muscular dystrophy (FCMD), mRNA
NM_006731	Homo sapiens deoxyribonuclease I-like 1 (DNASE1L1), mRNA
NM_006730	
NM_004366	
NM_006725	1 2 (D(Y,Y) 122 P(N))
NM_006806	The DTC family member 2 (RT(12) MKNA
NM_006763	
NM_006789	
NM 006793	Homo sapiens peroxiredoxin 3 (PRDX3), nuclear gene encoding mitochondia
1	mPNA
NM 006818	TXALL1 fixed gone from chromosome 1g (AFIQ), mRNA
NM 004289	

NM 006526	Homo sapiens zinc finger protein 217 (ZNF217), mRNA
NM_006523	Homo sapiens X-prolyl aminopeptidase (aminopeptidase P)-like (XPNPEPL), mRNA
NM 006537	Homo sapiens ubiquitin specific protease 3 (USP3), mRNA
NM 006564	Homo sapiens G protein-coupled receptor (TYMSTR), mRNA
NM 006573	Homo sapiens tumor necrosis factor (ligand) superfamily, member 13b
	(TNFSF13B), mRNA
NM_001561	Homo sapiens tumor necrosis factor receptor superfamily, member 9
	(TNFRSF9), mRNA
NM 006528	Homo sapiens tissue factor pathway inhibitor 2 (TFPI2), mRNA
NM 006520	Homo sapiens t-complex-associated-testis-expressed 1-like (TCTE1L), mRNA
NM 006519	Homo sapiens t-complex-associated-testis-expressed 1-like 1 (TCTEL1), mRNA
NM_006602	Homo sapiens transcription factor-like 5 (basic helix-loop-helix) (TCFL5),
11111_000002	mRNA
NM_006593	Homo sapiens T-box, brain, 1 (TBR1), mRNA
NM_006679	Homo sapiens putative opioid receptor, neuromedin K (neurokinin B) receptor-
14141_000073	like (TAC3RL), mRNA
NM 006682	Homo sapiens fibrinogen-like 2 (FGL2), mRNA
NM 006558	Homo sapiens Sam68-like phosphotyrosine protein, T-STAR (T-STAR), mRNA
NM 006603	Homo sapiens stromal antigen 2 (STAG2), mRNA
NM 006717	Homo sapiens spindlin (SPIN), mRNA
NM 006542	Homo sapiens S-phase response (cyclin-related) (SPHAR), mRNA
NM 00654	Homo sapiens suc1-associated neurotrophic factor target (FGFR signalling
19191_000034	adaptor) (SNT-1), mRNA
ND4 006622	Homo sapiens serum-inducible kinase (SNK), mRNA
NM_006622	Homo sapiens thyroid hormone receptor coactivating protein (SMAP), mRNA
NM_006696	Homo sapiens solute carrier family 2 (facilitated glucose transporter), member 1
NM_006516	(SLC2A1), mRNA
ND4 006622	Homo sapiens solute carrier family 17 (sodium phosphate), member 3
NM_006632	(SLC17A3), mRNA
ND 4 00 (517	Homo sapiens solute carrier family 16 (monocarboxylic acid transporters),
NM_006517	member 2 (putative transporter) (SLC16A2), mRNA
ND ( 006508	Homo sapiens solute carrier family 12 (potassium/chloride transporters), member
NM_006598	Homo sapiens solute carrier failing 12 (potassium emorite transporters), memoer
NB 6 00 6515	7 (SLC12A7), mRNA  Homo sapiens SET domain and mariner transposase fusion gene (SETMAR),
NM_006515	
) D. C. 006664	mRNA Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 27
NM_006664	
ND 6 006514	(SCYA27), mRNA  Homo sapiens sodium channel, voltage-gated, type X, alpha polypeptide
NM_006514	
377 6 006550	(SCN10A), mRNA  Homo sapiens GAP-associated tyrosine phosphoprotein p62 (Sam68) (SAM68),
NM_006559	
277 6006511	mRNA Homo sapiens regulatory solute carrier protein, family 1, member 1 (RSC1A1),
NM_006511	
277.6.006500	mRNA
NM_006583	Homo sapiens retinal pigment epithelium-derived rhodopsin homolog (RRH),
277 600000	mRNA
NM_006604	Homo sapiens ret finger protein-like 3 (RFPL3), mRNA
NM_006605	Homo sapiens ret finger protein-like 2 (RFPL2), mRNA
NM_006505	Homo sapiens poliovirus receptor (PVR), mRNA
NM_006504	Homo sapiens protein tyrosine phosphatase, receptor type, E (PTPRE), mRNA
NM_006503	Homo sapiens proteasome (prosome, macropain) 26S subunit, ATPase, 4
	(PSMC4), mRNA

WO 03/074654

WO 03/074654	
	Homo sapiens corin (PRSC), mRNA
	Homo sapiens corin (PRSC), mRNA  Homo sapiens phosphomevalonate kinase (PMVK), mRNA  Homo sapiens phosphomevalonate kinase (PMVK), mRNA
NM_006556	Homo sapiens phosphomevalonate kinase (PMVK), hikivit Homo sapiens putative homeodomain transcription factor (PHTF1), mRNA
NM_006608	
NM_006661	Homo sapiens phosphodiesterase 1912 (P5-1), mRNA  Homo sapiens MHC class I region ORF (P5-1), mRNA  to family 5, subfamily I, member 1 (OR5II),
NM_006674	Homo sapiens MHC class I region ORF (P3-1), mixty.  Homo sapiens olfactory receptor, family 5, subfamily I, member 1 (OR511),
NM_006637	Homo sapiens offactory 2007
70.5540	mRNA Homo sapiens serologically defined colon cancer antigen 16 (SDCCAG16),
NM_006649	
100500	1
NM_002532	Homo sapiens nucleoporin sort (Nervey).  Homo sapiens neuropathy target esterase (NTE), mRNA  Homo sapiens neuropathy target esterase (NTE), mRNA  sapiens nucleoporin sort (NEVEY).
NM_006702	Homo sapiens neuropathy target esterase (NTE), mid VI Homo sapiens cleavage and polyadenylation specific factor 4, 30kD subunit
NM_006693	(CDSE4) mRNA
	(CPSF4), mRNA Homo sapiens leukocyte immunoglobulin-like receptor, subfamily B (with TM
NM_006669	Homo sapiens leukocyte mindategichen and ITIM domains), member 1 (LILRB1), mRNA and ITIM domains), member 1 (LILRB1), mRNA
	and ITIM domains), member 1 (Enrice 1), and ITIM domains), member 1 (Enrice 2), and ITIM domains), member 1 (Enrice 2), and ITIM domains), member 1 (Enrice 2), and and ITIM domains), member 2 (Enrice 2), and and Enrice 2), and and Enrice 2 (Enrice 2), and and Enrice 2 (Enri
NM_006533	Homo sapiens melanoma inhibitory activity (MASS), and Homo sapiens melanoma adhesion molecule (MCAM), mRNA  Homo sapiens melanoma adhesion molecule (MCAM), mRNA  Linding lectin serine protease 2 (MASP2), mRNA
NM_006500	Homo sapiens melanoma adhesion molecule (MC/MX), MRNA  Homo sapiens mannan-binding lectin serine protease 2 (MASP2), mRNA  Homo sapiens mannan-binding lectin serine protease 2 (MAN1A2), mRNA
NM_006610	Homo sapiens mannan-binding lectin serine protease 2 (MAN1A2), mRNA  Homo sapiens mannosidase, alpha, class 1A, member 2 (MAN1A2), mRNA  Homo sapiens mannosidase, alpha, class 1A, member 2 (MAN1A2), mRNA  Homo sapiens mannosidase, alpha, class 1A, member 2 (MAN1A2), mRNA  Homo sapiens mannan-binding lectin serine protease 2 (MAN1A2), mRNA  Homo sapiens mannan-binding lectin serine protease 2 (MAN1A2), mRNA
NM_006699	Homo sapiens mannosidase, alpha, class 1A, memocr 2 (magazina), Homo sapiens lectin, galactoside-binding, soluble, 2 (galectin 2) (LGALS2),
NM_006498	Homo sapiens reem, gazare
	mRNA Homo sapiens IGF-II mRNA-binding protein 3 (KOC1), mRNA  Homo sapiens IGF-II mRNA-binding protein 3 (KOC1), mRNA  (KLRA1),
NM_006547	Homo sapiens IGF-II mRNA-binding protein 3 (KOC1), interest 1 (KLRA1), Homo sapiens killer cell lectin-like receptor subfamily A, member 1 (KLRA1),
NM_006611	Homo sapiciis kiner con activity
	mRNA Homo sapiens IGF-II mRNA-binding protein 1 (IMP-1), mRNA  GREEN mRNA
NM_006546	Homo sapiens heparanase (HPSE), mRNA  Homo sapiens heparanase (HPSE), mRNA
NM_006665	
NM_006497	Homo sapiens hypermetnylated in career (HERC2), mRNA  Homo sapiens heet domain and RLD 2 (HERC2), mRNA
NM_004667	Transport Hairpin hinding blotchi, mistorio (1-1-7)
NM_006527	Homo sapiens G-substrate (GSBS), mRNA  Homo sapiens G-substrate (GSBS), mRNA  placetide hinding protein (G protein), alpha inhibiting
NM_006658	
NM_006496	
NM_006529	Homo sapiens glycine receptor, aipha 5 (GES41), mRNA Homo sapiens glioma-amplified sequence-41 (GAS41), mRNA (GES41), mRNA (GES41), mRNA (GES41), mRNA
NM_00653	
NM_00658	1 Homo sapiens lucosyltanisterae
	mRNA 0 Homo sapiens FLN29 gene product (FLN29), mRNA 0 Homo sapiens FLN29 gene product (FLN29), mRNA
NM_00670	Homo sapiens FLN29 gene product (TEN25), mRNA Homo sapiens complement factor H-related 4 (FHR-4), mRNA  Homo sapiens FLN29 gene product (TEN25), mRNA
NM_00668	
NM_00411	Homo sapiens fibroblast growth factor 122 (EVI2B), mRNA Homo sapiens ecotropic viral integration site 2B (EVI2B), mRNA  (11.10 bysine rich leukemia gene) (ELL), mRNA
NM_00649	
NM_00653	Homo sapiens adhesion glycoprotein (DNAM-1), mRNA Homo sapiens adhesion glycoprotein (CYSLT1), mRNA
NM_00656	
NM_0066	Homo sapiens cysteinyl leukotrielle receptor (TNRC5), mRNA  Homo sapiens trinucleotide repeat containing 5 (TNRC5), mRNA  Liver factor (zinc finger protein) (CTCF), mRNA
NM_0065	Homo sapiens trinucleotide repeat containing 5 (Trices), including 1866 Homo sapiens CCCTC-binding factor (zinc finger protein) (CTCF), mRNA Homo sapiens CCCTC-binding factor (zinc finger protein) (CTCF), mRNA  (CSPG5),
NM_0065	
NM_0065	74 Homo sapiens chondrollin sunate protesty
	mRNA (CRE) mRNA
NM_0066	
NM_0064	Homo sapiens ceroid-inpoluseinosis, ileas
NM 0017	Homo sapiens calpastatin (CAS1), mile (1)
NM_0066	
NM_0066	

),
2),
),
),
), ),
),
),

1	nRNA (451 D) (SDE45) mRNA
M 006450	Homo sapiens splicing factor (45kD) (SPF45), mRNA
JM_006422	Homo sapiens splicing factor (45RD) (S1T-45), interest (45RD), mRNA Homo sapiens A kinase (PRKA) anchor protein 3 (AKAP3), mRNA  (AKAP3), mRNA
JM 006446	Homo sapiens solute carrier family 21 (organic amon acceptance)
VM_006278	(SLC21A6), mRNA  Homo sapiens sialyltransferase 4C (beta-galactosidase alpha-2,3-
NM_006378	1
[NIVI_000570	Homo sapiens sema domain, iminunogioodim domain (ES), domain (TM) and short cytoplasmic domain, (semaphorin) 4D (SEMA4D),
NM_006379	Homo saniens sema domain, immunoglobulin domain (1g), shore base
NN_000373	
NM_006274	Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 19
14141_000274	(GOVA 10) mPNA
NM 006453	1 (hata) like 3 (URL 1) MKNA
NM_006270	II and geniens retinitis nigmentosa I (autosolilai dolliman) (22 - 72
NM_006269	Homo gapiens ring finger protein 15 (RNF13), IIIRIVA
NM_006355	Homo sapiens ring finger protein 3 (RNF3), mRNA  Homo sapiens ring finger protein 3 (RNF3), mRNA
NM_006315	
NM_006394	Homo sapiens regulated in ghoma (RiO), interview Homo sapiens proteasome (prosome, macropain) activator subunit 1 (PA28)
NM_006263	Homo sapiens proteasonie (prosonie, maerepass)
	alpha) (PSME1), mRNA
NM_006262	Homo sapiens peripherin (PRPH), mRNA  Homo sapiens prophet of Pit1, paired-like homeodomain transcription factor
NM 006261	Homo sapiens prophet of Pitt, palled-like homeoderna
	(PROP1), mRNA  Homo sapiens protein-kinase, interferon-inducible double stranded RNA
NM 006260	Homo sapiens protein-kinase, interferon-finduciore double de la
NM_006259	Homo sapiens protein kinase, cGMP-dependent, type II (PRKG2), mRNA
NM 006257	III and comient protein kinase C. Inela (PRACQ), mattr
NM 006255	
NM_006253	Homo sapiens protein kinase C, eta (FRRCII), med War Homo sapiens protein kinase, AMP-activated, beta 1 non-catalytic subunit
14141_000200	
NM_006252	(PRKAB1), mRNA  Homo sapiens protein kinase, AMP-activated, alpha 2 catalytic subunit
14141_000252	
NM_006251	(PRKAA2), mRNA  Homo sapiens protein kinase, AMP-activated, alpha 1 catalytic subunit
1411_000251	1 · · · · · · · · · · · · · · ·
NM_006247	- In ambotoce 5 CatalVIIC SUDUIII (11115C), 1111C II
NM_006246	Homo sapiens protein phosphatase 2, regulatory subunit B (B56), epsilon
NW_000240	isoform (PPP2R5E), mRNA
NM_006245	
NM_000243	(PPP2R5D), mRNA
27.5.00(244	- 1- ambatace / regulatory subulit D (D30), out 120
NM_006244	
NM_006243	
	(PPP2R5A), mRNA  Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 2 (PPP1R2)
NM_006241	
	mRNA Homo sapiens protein phosphatase, EF hand calcium-binding domain 1 (PPEF1
NM_006240	
	mRNA  Homo sapiens peroxisome proliferative activated receptor, delta (PPARD),
NM_00623	
	mRNA Homo sapiens POU domain, class 4, transcription factor 1 (POU4F1), mRNA
NM_00623	Homo sapiens POU domain, class 4, transcription ractor 1 (2 2 2 2)

35.0000	TI
NM_006236	Homo sapiens POU domain, class 3, transcription factor 3 (POU3F3), mRNA Homo sapiens POU domain, class 2, associating factor 1 (POU2AF1), mRNA
NM_006235	Homo sapiens POU domain, class 2, associating factor 1 (1002Ar1), filed 1.  Homo sapiens polymerase (DNA directed), epsilon (POLE), mRNA
NM_006231	Homo sapiens polymerase (DNA directed), epsilon (1 OLD), index.  Homo sapiens solute carrier family 25 (mitochondrial carrier; peroxisomal
NM_006358	Homo sapiens solute carrier family 25 (filliconomicial carrier, peroxisornal
	membrane protein, 34kD), member 17 (SLC25A17), mRNA
NM_006227	Homo sapiens phospholipid transfer protein (PLTP), mRNA
NM_006226	Homo sapiens phospholipase C, epsilon (PLCE), mRNA
NM_006225	Homo sapiens phospholipase C, delta 1 (PLCD1), mRNA
NM_006224	Homo sapiens phosphotidylinositol transfer protein (PITPN), mRNA
NM_006479	Homo sapiens RAD51-interacting protein (PIR51), mRNA
NM_006223	Homo sapiens protein (peptidyl-prolyl cis/trans isomerase) NIMA-interacting, 4 (parvulin) (PIN4), mRNA
NM 006222	Homo sapiens protein (peptidyl-prolyl cis/trans isomerase) NIMA-interacting 1-
19191_000222	like (PIN1L), mRNA
NM 006221	Homo sapiens protein (peptidyl-prolyl cis/trans isomerase) NIMA-interacting 1
NWI_000221	(PIN1), mRNA
NM 006218	Homo sapiens phosphoinositide-3-kinase, catalytic, alpha polypeptide
1 - 1 - 1	(PIK3CA), mRNA
NM 006213	Homo sapiens phosphorylase kinase, gamma 1 (muscle) (PHKG1), mRNA
NM 006305	Homo sapiens putative human HLA class II associated protein I (PHAP1),
_	mRNA
NM 006212	Homo sapiens 6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 2
_	(PFKFB2), mRNA
NM 006211	Homo sapiens proenkephalin (PENK), mRNA
NM 006209	Homo sapiens ectonucleotide pyrophosphatase/phosphodiesterase 2 (autotaxin)
_	(ENPP2), mRNA
NM_006205	Homo sapiens phosphodiesterase 6H, cGMP-specific, cone, gamma (PDE6H), mRNA
NM 006204	Homo sapiens phosphodiesterase 6C, cGMP-specific, cone, alpha prime
	(PDE6C), mRNA
NM 006198	Homo sapiens Purkinje cell protein 4 (PCP4), mRNA
NM 006197	Homo sapiens pericentriolar material 1 (PCM1), mRNA
NM 006195	Homo sapiens pre-B-cell leukemia transcription factor 3 (PBX3), mRNA
NM 006193	Homo sapiens paired box gene 4 (PAX4), mRNA
NM 006191	Homo sapiens proliferation-associated 2G4, 38kD (PA2G4), mRNA
NM 006189	Homo sapiens olfactory marker protein (OMP), mRNA
NM 006186	Homo sapiens nuclear receptor subfamily 4, group A, member 2 (NR4A2),
	mRNA
NM 006185	Homo sapiens nuclear mitotic apparatus protein 1 (NUMA1), mRNA
NM 006184	Homo sapiens nucleobindin 1 (NUCB1), mRNA
NM 006182	Homo sapiens discoidin domain receptor family, member 2 (DDR2), mRNA
NM 006180	Homo sapiens neurotrophic tyrosine kinase, receptor, type 2 (NTRK2), mRNA
NM 006372	Homo sapiens NS1-associated protein 1 (NSAP1), mRNA
NM 006177	Homo sapiens neural retina leucine zipper (NRL), mRNA
NM 006176	Homo sapiens neurogranin (protein kinase C substrate, RC3) (NRGN), mRNA
NM 006174	Homo sapiens neuropeptide Y receptor Y5 (NPY5R), mRNA
NM 006170	Homo sapiens nucleolar protein 1 (120kD) (NOL1), mRNA
NM 006169	Homo sapiens nicotinamide N-methyltransferase (NNMT), mRNA
NM 006165	Homo sapiens nuclear factor related to kappa B binding protein (NFRKB),
1111_000103	mRNA
NM_006164	Homo sapiens nuclear factor (erythroid-derived 2)-like 2 (NFE2L2), mRNA
NM 006163	Homo sapiens nuclear factor (erythroid-derived 2), 45kD (NFE2), mRNA
14141 000103	1 20000 100000 10000 (0) 10000 - 10000 (0)

	Homo sapiens neurogenic differentiation 2 (NEUROD2), mRNA
NM_006160 ]	Homo sapiens neurogeme differentiation 2 (NZESTA), mRNA Homo sapiens neurofilament, light polypeptide (68kD) (NEFL), mRNA
	Homo sapiens neuromament, fight polypeptide (5512)
	Homo sapiens nebulette (NEBL), mRNA Homo sapiens DNA-binding transcriptional activator (NCYM), mRNA
	Homo sapiens DNA-binding transcriptional detributes (**)
	Homo sapiens NCK adaptor protein 1 (NCK1), mRNA Homo sapiens solute carrier family 34 (sodium phosphate), member 2
NM_006424	Homo sapiens solute carrier failily 34 (sounding prosperator),
	(SLC34A2), mRNA Homo sapiens brain acid-soluble protein 1 (BASP1), mRNA
NM_006317	Homo sapiens brain acid-soluble protein 1 (Brish 1), mRNA  Homo sapiens c-mer proto-oncogene tyrosine kinase (MERTK), mRNA
	Homo sapiens c-mer proto-oncogene tyrosine kinase (***) Homo sapiens LIM protein (similar to rat protein kinase C-binding enigma)
NM_006457	(LIM) mRNA
NM 006148	T TM and SU2 protein 1 (LASPI), MKNA
	Homo sapiens DNA-dependent protein kinase catalytic subunit-interacting
NM_006383	· • azmo\DNIA
27.5.006450	Homo sapiens similar to Caenorhabditis elegans protein C42C1.9 (KEO4),
NM_006459	DATA
ND 6 000147	mRNA  Homo sapiens interferon regulatory factor 6 (IRF6), mRNA  Output
NM_006147	Homo conjens interferon gamma-inducible protein 30 (H-130), find (12)
NM_006332	Homo sapiens microspherule protein 1 (MCRS1), mRNA  Homo sapiens microspherule protein 1 (MCRS1), mRNA
NM_006337	1 4 -1 ole 27kl) protein 1 (HSPB2), IIINNA
NM_006308	Homo sapiens heat snock 27kD protein 3 (116125),  Homo sapiens enhancer of filamentation 1 (cas-like docking; Crk-associated
NM_006403	Homo sapiens ennancer of mathemation 1 (our first of the sapiens ennancer of mathematical first of the sapiens ennancer of the sapiens ennan
	substrate related) (HEF1), mRNA  Homo sapiens G protein-coupled receptor 19 (GPR19), mRNA
NM_006143	Homo sapiens G protein-coupled receptor 15 (GTEC),
NM_006302	Homo sapiens glucosidase I (GCS1), mRNA  Homo sapiens glucosidase I (GCS1), mRNA
NM_006478	Homo sapiens GAS2-related on chromosome 22 (GAR22), mRNA  Homo sapiens GAS2-related on chromosome 1 protein (leucine-rich)
NM_006338	Homo sapiens GAS2-related on chromosome 1 protein (leucine-rich)
	(GAC1), mRNA
NM_006360	Homo sapiens dendritic cell protein (GA17), mRNA
NM_006329	Homo sapiens fibulin 5 (FBLN5), mRNA  Homo sapiens fibulin 5 (FBLN5), mRNA
NM_006404	Homo sapiens frouin 5 (FBEN5), find 11  Homo sapiens protein C receptor, endothelial (EPCR) (PROCR), mRNA  Homo sapiens protein C receptor, endothelial (EPCR) (PROCR), mRNA
NM 006304	Homo sapiens Deleted in split-hand/split-foot 1 region (DSS1), mRNA  Homo sapiens Deleted in split-hand/split-foot 1 region (DSS1), mRNA
NM 001355	Homo sapiens D-dopachrome tautomerase (DDT), mRNA
NM 006139	Homo sapiens CD28 antigen (Tp44) (CD28), mRNA  Homo sapiens CD28 antigen (Tp44) (CD28), mRNA
NM 006371	Training aggregated protein (CR IAF), IIINNA
NM_006136	Homo sapiens cartilage associated protein (example 2) Homo sapiens capping protein (actin filament) muscle Z-line, alpha 2
	(CADZAO)DNA
NM 006448	Homo sapiens trinucleotide repeat containing 1 (TNRC1), mRNA
NM 006333	$1_{-}$ $1_{-$
NM 006419	Homo saniens small inducible cytokine B subfamily (Cys-A-Cys motif), memora
	12 (P. sell chemoattractant) (SCYBI3), mRNA
NM 005453	TT since since finger protein 79/(ZNF29/), IIIKNA
NM 006324	development protein 1 (CFD1 1), interes
NM 006375	Homo sapiens cytosolic ovarian carcinoma antigen I (COVAI), mid-VI
NM_004466	Homo sapiens glypican 5 (GPC5), mRNA
NM_004484	· 1 ··································
NM 002856	Homo sapiens glypican 3 (GrC3), interview Homo sapiens poliovirus receptor-related 2 (herpesvirus entry mediator B)
14141_005020	
NM 001420	(PVRL2), mRNA  Homo sapiens ELAV (embryonic lethal, abnormal vision, Drosophila)-like 3 (H
14141_001420	
NM 001634	Homo saniens S-adenosylmethionine decarboxylase I (AWDI), Illication
19191_001034	
NM 000483	Homo sapiens apolipoprotein C-I (APOC1), mRNA

NM_000482	Homo sapiens apolipoprotein A-IV (APOA4), mRNA
NM_005953	Homo sapiens metallothionein 2A (MT2A), mRNA
NM_005954	Homo sapiens metallothionein 3 (growth inhibitory factor (neurotrophic))
	(MT3), mRNA
NM_006007	Homo sapiens zinc finger protein 216 (ZNF216), mRNA
NM_006006	Homo sapiens zinc finger protein 145 (Kruppel-like, expressed in promyelocytic
	leukemia) (ZNF145), mRNA
NM_006004	Homo sapiens ubiquinol-cytochrome c reductase hinge protein (UQCRH),
	mRNA
NM_006003	Homo sapiens ubiquinol-cytochrome c reductase, Rieske iron-sulfur polypeptide
	1 (UQCRFS1), nuclear gene encoding mitochondrial protein, mRNA
NM_006088	Homo sapiens tubulin, beta, 2 (TUBB2), mRNA
NM_005999	Homo sapiens translin-associated factor X (TSNAX), mRNA
NM_006022	Homo sapiens transforming growth factor beta-stimulated protein TSC-22
) D. f. 005000	(TSC22), mRNA
NM_005998	Homo sapiens chaperonin containing TCP1, subunit 3 (gamma) (CCT3), mRNA
NM_006073	Homo sapiens triadin (TRDN), mRNA
NM_005997	Homo sapiens transcription factor-like 1 (TCFL1), mRNA
NM_006116	Homo sapiens transforming growth factor beta-activated kinase-binding protein 1 (TAB1), mRNA
NM 005989	Homo sapiens aldo-keto reductase family 1, member D1 (delta 4-3-ketosteroid-
INIVI_003989	5-beta-reductase) (AKR1D1), mRNA
NM 005988	Homo sapiens small proline-rich protein 2A (SPRR2A), mRNA
NM 005986	Homo sapiens SRY (sex determining region Y)-box 1 (SOX1), mRNA
NM_006049	Homo sapiens small nuclear RNA activating complex, polypeptide 5, 19kD
14141_000049	(SNAPC5), mRNA
NM 006080	Homo sapiens sema domain, immunoglobulin domain (Ig), short basic domain,
_	secreted, (semaphorin) 3A (SEMA3A), mRNA
NM_006072	Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 26
	(SCYA26), mRNA
NM_005981	Homo sapiens sarcoma amplified sequence (SAS), mRNA
NM_006054	Homo sapiens reticulon 3 (RTN3), mRNA
NM_005977	Homo sapiens ring finger protein (C3H2C3 type) 6 (RNF6), mRNA
NM_005975	Homo sapiens PTK6 protein tyrosine kinase 6 (PTK6), mRNA
NM_005972	Homo sapiens pancreatic polypeptide receptor 1 (PPYR1), mRNA
NM_006112	Homo sapiens peptidylprolyl isomerase E (cyclophilin E) (PPIE), mRNA
NM_006107	Homo sapiens acid-inducible phosphoprotein (OA48-18), mRNA
NM_006067	Homo sapiens neighbor of COX4 (NOC4), mRNA
NM_005969	Homo sapiens nucleosome assembly protein 1-like 4 (NAP1L4), mRNA
NM_006058	Homo sapiens Nef-associated factor 1 (NAF1), mRNA
NM_006097	Homo sapiens myosin regulatory light chain 2, smooth muscle isoform
ND 4 005055	(MYRL2), mRNA
NM_005955	Homo sapiens metal-regulatory transcription factor 1 (MTF1), mRNA
NM_005932	Homo sapiens mitochondrial intermediate peptidase (MIPEP), nuclear gene
NIM 005021	encoding mitochondrial protein, mRNA  Home serious MIC class I relymentide related assumes B (MCR) RNA
NM_005931	Homo sapiens MHC class I polypeptide-related sequence B (MICB), mRNA
NM_006081 NM_005930	Homo sapiens MHC binding factor, beta (MHCBFB), mRNA  Homo sapiens maniprisms expressed antigen 6 (acided acid maline rich)
14141 002320	Homo sapiens meningioma expressed antigen 6 (coiled-coil proline-rich) (MGEA6), mRNA
NM 005928	Homo sapiens milk fat globule-EGF factor 8 protein (MFGE8), mRNA
NM 005926	Homo sapiens microfibrillar-associated protein 1 (MFAP1), mRNA
NM 005925	Homo sapiens meprin A, beta (MEP1B), mRNA
1111 000720	1 rome suprens meprin A, ocu (MDI ID), mulia

VM_005924	Homo sapiens mesenchyme homeo box 2 (growth arrest-specific homeo box)
	(MEOX2), mRNA  Homo sapiens MADS box transcription enhancer factor 2, polypeptide D  (MEEXD) mRNA
VM_005920	Homo sapiens MADS box transcription eminancer rates 2, poster 1
	(myocyte enhancer factor 2D) (MEF2D), mRNA  Homo sapiens MADS box transcription enhancer factor 2, polypeptide B
NM 005919	Homo sapiens MADS box transcription eminated research and the sapiens MADS box transcription eminated research
	(myocyte enhancer factor 2B) (MEF2B), mRNA  Homo sapiens malate dehydrogenase 2, NAD (mitochondrial) (MDH2), nuclear
NM_005918	Homo sapiens malate dehydrogenase 2, 1711 (miles)
	gene encoding mitochondrial protein, mRNA  Homo sapiens malate dehydrogenase 1, NAD (soluble) (MDH1), mRNA  Homo sapiens malate dehydrogenase 1, NAD (soluble) (MDH1), mRNA
NM 005917	Homo sapiens malate denydrogenase 1, 1471 (6518-7),
NM 005913	
NM 005912	
NM 005911	Homo sapiens methionine adenosyllatisticase in, approximation of the sapiens methionine adenosyllatisticase in the sapiens methion of the sapiens method methion of the sapiens methion of the sapiens methion of the sapiens methion of the sapiens method met
NM 005908	Homo sapiens metnionine adenosyntanoreces (MANBA), mRNA Homo sapiens mannosidase, beta A, lysosomal (MANBA), mRNA
NM 005907	
NM_005898	Homo sapiens membrane component, en omosomo 23,
11112	(M11S1), mRNA
NM 006060	II was senious zinc finger protein, sublamily IA, I (Ikaros) (21,121,197)
NM 006059	
NM 006038	
NM_006084	Homo sapiens spermatogenesis associated 151 (122 22 22 22 22 22 22 22 22 22 22 22 22
14141_00000-1	(ISGF3G), mRNA
NM 005897	
NM 005896	
NM 006028	
NM_006120	Homo sapiens 5-hydroxytryptamine (serotomi) receptation of the Homo sapiens major histocompatibility complex, class II, DM alpha (HLA-Homo sapiens major histocompatibility complex, class II, DM alpha (HLA-Homo sapiens major histocompatibility complex).
NM_000120	- x s A \ TD \ X A
NM 006026	There's ganiene H1 histone family, member X (H1FA), HIXIVA
NM 006020	
NM_006031	Homo saniens Chp/p300-interacting transactivator, with
NM_006079	terminal domain, 2 (CITED2), mRNA
274 005904	terminal domain, 2 (CITED2), mRNA  Homo sapiens CD5 antigen-like (scavenger receptor cysteine rich family)
NM_005894	
ND 6 000016	and continue giolomycin (CI) (D4), IIIXIXA
NM_006016	
NM_006078	(CACNG2), mRNA
5 5 00 6020	
NM_006030	(CACNA2D2), mRNA
NM_006085	Transport CWIT/SNE related, matrix associated, admir deposition
NM_006015	
NM_00606	(AKR1A1), mRNA
	1 Common A acetyltransferase 2 (acetylectyl Cooling) into
NM_00589	1 1 1 1 1 (A('A'I')) mRNA
	There agains alloylation repair; alkB homolog (ABII), inclus
NM_00602	- tare combonic onbydrase VIII (CAO), IIICVA
NM 00405	- 1 mag tinger projetti. J (IVII)
NM_00566	1/ - Jamondent anion (III/IIIIC) J ( 1 D1100 /)
NM_00566	Homo sapiens voltage-dependent amon chamber 2  Homo sapiens translational inhibitor protein p14.5 (UK114), mRNA  Homo sapiens translational inhibitor protein p14.5 (UK114), mRNA
NM_00583	
NM_00566	
	(SLC35A2), mRNA Homo sapiens ubiquitin fusion degradation 1-like (UFD1L), mRNA

NM 005706	Homo sapiens tumor suppressing subtransferable candidate 4 (TSSC4), mRNA
NM 005723	Homo sapiens tetraspan 5 (TSPAN-5), mRNA
NM 005727	Homo sapiens tetraspan 1 (TSPAN-1), mRNA
NM 005658	Homo sapiens TNF receptor-associated factor 1 (TRAF1), mRNA
	Homo sapiens tumor protein p53-binding protein (TP53BPL), mRNA
NM_005802	Homo sapiens transducer of ERBB2, 1 (TOB1), mRNA
NM_005749	Homo sapiens TGFB inducible early growth response (TIEG), mRNA
NM_005655	Homo sapiens transcription factor CP2 (TFCP2), mRNA
NM_005653	Homo sapiens nuclear receptor subfamily 2, group F, member 1 (NR2F1),
NM_005654	mRNA
NM_005652	Homo sapiens telomeric repeat binding factor 2 (TERF2), mRNA
NM 005885	Homo sapiens similar to S. cerevisiae SSM4 (TEB4), mRNA
NM 005651	Homo sapiens tryptophan 2,3-dioxygenase (TDO2), mRNA
NM 005649	Homo sapiens transcription factor 17 (TCF17), mRNA
NM 005647	Homo sapiens transducin (beta)-like 1 (TBL1), mRNA
NM_005645	Homo sapiens TATA box binding protein (TBP)-associated factor, RNA
	polymerase II, K, 18kD (TAF2K), mRNA
NM_005643	Homo sapiens TATA box binding protein (TBP)-associated factor, RNA polymerase II, I, 28kD (TAF2I), mRNA
NTM 005641	Homo sapiens TATA box binding protein (TBP)-associated factor, RNA
NM_005641	polymerase II, E, 70/85kD (TAF2E), mRNA
NM 005679	Homo sapiens TATA box binding protein (TBP)-associated factor, RNA
14141_003079	polymerase I, C, 110kD (TAF1C), mRNA
NM 005681	Homo sapiens TATA box binding protein (TBP)-associated factor, RNA
MM_002091	polymerase I, A, 48kD (TAF1A), mRNA
NM 005639	Homo sapiens synaptotagmin 1 (SYT1), mRNA
NM 005638	Homo sapiens synaptotrevin-like 1 (SYBL1), mRNA
	Homo sapiens synovial sarcoma, X breakpoint 1 (SSX1), mRNA
NM 005635 NM 005871	Homo sapiens splicing factor 30, survival of motor neuron-related (SPF30),
14141_003671	mRNA
NM 005634	Homo sapiens SRY (sex determining region Y)-box 3 (SOX3), mRNA
NM 005686	Homo sapiens SRY (sex determining region Y)-box 13 (SOX13), mRNA
NM_005629	Homo sapiens solute carrier family 6 (neurotransmitter transporter, creatine),
	member 8 (SLC6A8), mRNA
NM_005630	Homo sapiens solute carrier family 21 (prostaglandin transporter), member 2 (SLC21A2), mRNA
NM 005628	Homo sapiens solute carrier family 1 (neutral amino acid transporter), member 5
1117_005020	(SLC1A5), mRNA
NM 005627	Homo sapiens serum/glucocorticoid regulated kinase (SGK), mRNA
NM 005877	Homo sapiens splicing factor 3a, subunit 1, 120kD (SF3A1), mRNA
NM 005625	Homo sapiens syndecan binding protein (syntenin) (SDCBP), mRNA
NM 005623	Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 8
1414_005025	(monocyte chemotactic protein 2) (SCYA8), mRNA
NM_005624	Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 25
1111_00002	(SCYA25), mRNA
NM_005850	Homo sapiens splicing factor 3b, subunit 4, 49kD (SF3B4), mRNA
NM_005772	Homo sapiens RNA cyclase homolog (RNAC), mRNA
NM 005614	The state of the s
	Homo sapiens Ras homolog enriched in brain 2 (RHEB2), mRNA
NM_005777	Homo sapiens RNA binding motif protein 6 (RBM6), mRNA
	Homo sapiens RNA binding motif protein 6 (RBM6), mRNA Homo sapiens RNA binding motif protein 5 (RBM5), mRNA
NM_005777	Homo sapiens RNA binding motif protein 6 (RBM6), mRNA

WO 03/074654

NM 005607	Homo sapiens PTK2 protein tyrosine kinase 2 (PTK2), mRNA
TNA 005780	Homo sapiens proteasome (prosonie, maeropani)
NM 005865	Homo sapiens prostate stem cen antigen (x 5003), mRNA Homo sapiens protease, serine, 16 (thymus) (PRSS16), mRNA Homo sapiens protease, serine, 16 (thymus) (PRSS16), mRNA
NM 005729	Homo sapiens protease, serine, 16 (thythus) (TROSTO), mRNA Homo sapiens peptidylprolyl isomerase F (cyclophilin F) (PPIF), mRNA Homo sapiens peptidylprolyl isomerase F (cyclophilin F) (POU3F2), mRNA
NM 005604	
NM 005709	Homo sapiens POU dolliam, class 5, additional Homo sapiens PDZ-73 protein (PDZ-73/NY-CO-38), mRNA  Homo sapiens PDZ-73 protein (PDZ-73/NY-CO-38), mRNA
NM 005767	
NM 005835	Homo sapiens purmergic receptor (talmy 11 group 5) (Homo sapiens solute carrier family 17 (sodium phosphate), member 2
MM_002922	(SLC17A2), mRNA
NM_005793	Home saniens nucleoside diphosphate killase type o (militare type o (milit
MM_003773	enentosis alpha) (NM23-H0), mixiva
NM_005600	
NM 005599	
NM 005598	TT regions nescient helix 1000 nellx 1 (11111111); Alle
NM 005596	
NM_005390	Homo sapiens nuclear factor by (14 by), interpretation site 5 (EVI5), mRNA  Homo sapiens ecotropic viral integration site 5 (EVI5), mRNA
	Homo sapiens ecotropic viral integration site 3 (2 v2); and the Homo sapiens nascent-polypeptide-associated complex alpha polypeptide
NM_005594	OLA CIA) mDNIA
ND 5 005502	
NM_005593	Homo sapiens myogenic factor 5 (MTF3), find TY Homo sapiens muscle, skeletal, receptor tyrosine kinase (MUSK), mRNA Homo sapiens muscle, skeletal, receptor tyrosine kinase (MUSK), mRNA Homo sapiens myogenic factor 5 (MTF3), find TY Homo sapiens muscle, skeletal, receptor tyrosine kinase (MUSK), mRNA
NM_005592	Homo sapiens muscle, skeletal, receptor tyrosine kmase (Webszy) Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 4
NM_005845	(ABCC4), mRNA
77.6.005074	Till a regions leukocyte immunoglobulin-like leceptor, suotaminy
NM_005874	and ITIM domains), member 2 (LILRB2), mRNA  and ITIM domains), member 2 (LILRB2), mRNA
NA 005500	
NM_005588	TI conjent MAI)S hox iranscription childhed 2200 - 71
NM_005587	(myocyte enhancer factor 2A) (MEF2A), mRNA  (myocyte enhancer factor 2A) (MEF2A), mRNA  (myocyte enhancer factor 2A) (MEF2A), mRNA
27.6 005010	(myocyte enhancer factor 2A) (MEF2A), mRNA  Homo sapiens killer cell lectin-like receptor subfamily G, member 1 (KLRG1),
NM_005810	mRNA : 1 1-1) (LII) mRNA
ND 4 005591	mRNA Homo sapiens Lutheran blood group (Auberger b antigen included) (LU), mRNA Homo sapiens Lutheran blood group (Figure 1) preferred translocation partner in lipoma
NM_005581	Homo sapiens Lutheran blood group (Auberger o antigen increase)  Homo sapiens LIM domain-containing preferred translocation partner in lipoma
NM_005578	(TDD) mPNA
27.6.005.577	TX - remions linearotein Ln(a) (LPA), mRNA
NM_005577	TI are comione lycyl oxidase-like I (LUALI), illiciti
NM_005576	
NM_005573	
NM_005572	T TO ( )
NM_005568	Transfer of the one HM(t)( histori Datute (Lin 1), the
NM_005780	Translantate dehydrogenase A (LL)1111), mid 11
NM_005566	1: 1: 1 /oncogene /4117111 / 114 / 12 1
NM_005564	
NM_00555	Homo sapiens ladinin 1 (EEE 2), mRNA Homo sapiens keratin 7 (KRT7), mRNA
NM_00555	
NM_00555	
	(KRT16), mRNA  Homo sapiens keratin, cuticle, ultrahigh sulphur 1 (KRN1), mRNA  (KRT16), mRNA
NM_00555	
NM_00555	The state of the s
NM_00555	
NM_00555	
NM_00583	Homo sapiens potassium large conditionale curves subfamily M, beta member 2 (KCNMB2), mRNA
1 =	subtamily M, beta member 2 (KOTHID2),

ND4 005540	Homo sapiens potassium voltage-gated channel, shaker-related subfamily,
NM_005549	member 10 (KCNA10), mRNA
NM 005548	Homo sapiens lysyl-tRNA synthetase (KARS), mRNA
NM 005547	Homo sapiens involucrin (IVL), mRNA
NM_005545	Homo sapiens immunoglobulin superfamily containing leucine-rich repeat
1/101_002242	(ISLR), mRNA
NM 005853	Homo sapiens iroquois-class homeodomain protein (IRX-2A), mRNA
NM 005544	Homo sapiens insulin receptor substrate 1 (IRS1), mRNA
NM 005543	Homo sapiens insulin-like 3 (Leydig cell) (INSL3), mRNA
NM 005542	Homo sapiens insulin induced gene 1 (INSIG1), mRNA
NM 005541	Homo sapiens inositol polyphosphate-5-phosphatase, 145kD (INPP5D), mRNA
NM 005539	Homo sapiens inositol polyphosphate-5-phosphatase, 40kD (INPP5A), mRNA
NM 005537	Homo sapiens inhibitor of growth 1 family, member 1 (ING1), mRNA
NM 005535	Homo sapiens interleukin 12 receptor, beta 1 (IL12RB1), mRNA
NM 005532	Homo sapiens interferon, alpha-inducible protein 27 (IFI27), mRNA
NM 005531	Homo sapiens interferon, gamma-inducible protein 16 (IFI16), mRNA
NM 005530	Homo sapiens isocitrate dehydrogenase 3 (NAD+) alpha (IDH3A), mRNA
NM 005808	Homo sapiens HYA22 protein (HYA22), mRNA
NM 005528	Homo sapiens heat shock 40kD protein 2 (HSPF2), mRNA
NM 005526	Homo sapiens heat shock transcription factor 1 (HSF1), mRNA
NM 005525	Homo sapiens hydroxysteroid (11-beta) dehydrogenase 1 (HSD11B1), mRNA
NM 005522	Homo sapiens homeo box A1 (HOXA1), mRNA
NM 005521	Homo sapiens homeo box 11 (T-cell lymphoma 3-associated breakpoint)
	(HOX11), mRNA
NM 005518	Homo sapiens 3-hydroxy-3-methylglutaryl-Coenzyme A synthase 2
_	(mitochondrial) (HMGCS2), mRNA
NM 005515	Homo sapiens homeo box HB9 (HLXB9), mRNA
NM 005516	Homo sapiens major histocompatibility complex, class I, E (HLA-E), mRNA
NM_005712	Homo sapiens HERV-H LTR-associating 1 (HHLA1), mRNA
NM_005844	Homo sapiens PERB11 family member in MHC class I region (HCGIX), mRNA
NM_005513	Homo sapiens general transcription factor IIE, polypeptide 1 (alpha subunit,
	56kD) (GTF2E1), mRNA
NM_005683	Homo sapiens G protein-coupled receptor 55 (GPR55), mRNA
NM_005684	Homo sapiens G protein-coupled receptor 52 (GPR52), mRNA
NM_005512	Homo sapiens glycoprotein A repetitions predominant (GARP), mRNA
NM_005851	Homo sapiens tumor suppressor deleted in oral cancer-related 1 (DOC-1R), mRNA
NM 005740	Homo sapiens dynein, axonemal, light polypeptide 4 (DNAL4), mRNA
NM 005872	Homo sapiens breast carcinoma amplified sequence 2 (BCAS2), mRNA
NM 005671	Homo sapiens reproduction 8 (D8S2298E), mRNA
NM 005800	Homo sapiens highly charged protein (D13S106E), mRNA
NM 005752	Homo sapiens C-type (calcium dependent, carbohydrate-recognition domain)
	lectin, superfamily member 1 (cartilage-derived) (CLECSF1), mRNA
NM 005507	Homo sapiens cofilin 1 (non-muscle) (CFL1), mRNA
NM 005825	Homo sapiens RAS guanyl releasing protein 2 (calcium and DAG-regulated)
	(RASGRP2), mRNA
NM 005773	Homo sapiens zinc finger protein 256 (ZNF256), mRNA
NM 005774	Homo sapiens zinc finger protein 255 (ZNF255), mRNA
NM 005504	Homo sapiens branched chain aminotransferase 1, cytosolic (BCAT1), mRNA
NM 005738	Homo sapiens ADP-ribosylation factor-like 4 (ARL4), mRNA
	1 1 0 (0.41 D) (4 D D CO)
NM 005731	Homo sapiens actin related protein 2/3 complex, subunit 2 (34 kD) (ARPC2),

IM_005719	Homo sapiens actin related protein 2/3 complex, subunit 3 (21 kD) (ARPC3),
	to the molymoris COII like (AI CD), interest
= 5.005050	Homo saniens A kinase (PRKA) anchor protein o (1999)
NM 002023	Homo sapiens fibromodulin (FMOD), mRNA  Homo sapiens fibromodulin (FMOD), mRNA
NM_000108	debudrogenase complex, 2-0x0-glutarate complex, state-
	t t t t compley (III II) mking
	Homo sapiens aryl hydrocarbon receptor (AHR), mRNA
NM_001621	
NM_001101	
NM_001100	Homo sapiens actin, alpha 1, skeletal muscle (ACTAT), manual muscle
NM_000054	Homo sapiens arginine vasopressing and the same and the same arginine vasopressing argin
	(AVPR2), mRNA  Homo sapiens zinc finger protein 265 (ZNF265), mRNA  Homo sapiens zinc finger protein 265 (ZNF265), mRNA
NM_005455	Homo sapiens zinc finger protein 265 (ZNF265), mictri  Homo sapiens v-yes-1 Yamaguchi sarcoma viral oncogene homolog 1 (YES1),
NM_005433	Homo sapiens v-yes-1 Yamaguchi sarooma value
	mRNA Labelial growth factor C (VEGFC), mRNA
NM 005429	mRNA  Homo sapiens vascular endothelial growth factor C (VEGFC), mRNA  Homo sapiens vascular endothelial growth factor C (VEGFC), mRNA
NM 005499	Homo saniens SIIMO-1 activating enzyme subunit 2 (CD12),
NM 005427	
NM_005425	Homo sapiens tumor protein p/3 (1F/3), ind (1F/3), Homo sapiens transition protein 2 (during histone to protamine replacement)
14141_005 125	(TNP2), mRNA
NM_005424	Homo sapiens tyrosine kinase with immunoglobulin and epiderman grant
14141_003.2.	
NM_005423	Homo caniens trefoil factor 2 (spasmolytic protein 1) (1112),
NM 005422	
NM 005421	T call coute lymphocytic leukellila 2 (17xD2); x22
	Ifotroncterace estroycol-pictoring (012), 122
NM_005420	
NM_005418	domain binding protein 1 (Box13211)
NM_005470	Homo sapiens small proline-rich protein 3 (SPRR3), mRNA  Homo sapiens small proline-rich protein 3 (SPRR3), mRNA
NM_005416	
NM_005460	Homo sapiens synuciein, aipha interacting protein (5)-7  Homo sapiens serine hydroxymethyltransferase 2 (mitochondrial) (SHMT2),
NM_005412	
	mRNA Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 13
NM_005408	
	(SCYA13), mRNA  Homo sapiens v-ral simian leukemia viral oncogene homolog A (ras related)
NM_005402	Homo sapiens v-ral simian leukenna vital onogene
	(RALA), mRNA
NM 005397	Homo sapiens podocalyxin-like (PODXL), mRNA  Homo sapiens podocalyxin-like (PODXL), mRNA
NM 005395	
NM 005394	
NM_005390	
NM_005389	Homo sapiens protein-L-isoaspartate (D-aspartate) O-metry transcript
TVIVI_005505	(PCMT1) mRNA
NM 005450	Homo saniens noggin (NOG), mRNA
NM 00538	
NM_005384	1 footor interlenkin a regulated the interlenkin
	· 11 Jan 2 Confocult Signification INDOZI, mad 122
NM_00538	Homo saniens neurofilament 3 (150kD medium) (14L15); Medium
NM_00538	
NM_00538	
NM_00538	
NM_00546	Homo sapiens N-acetylated alpha-linked acidio dipopulated acidio dip
_	DIPEPTIDYLPEPTIDASE (NAALADAGEE), INCAT

NM_005374	Homo sapiens membrane protein, palmitoylated 2 (MAGUK p55 subfamily member 2) (MPP2), mRNA
NM 005373	Homo sapiens myeloproliferative leukemia virus oncogene (MPL), mRNA
NM_005373	Homo sapiens v-mos Moloney murine sarcoma viral oncogene homolog (MOS),
14141_005572	mRNA
NM 005439	Homo sapiens myeloid leukemia factor 2 (MLF2), mRNA
NM 005369	Homo sapiens MCF.2 cell line derived transforming sequence (MCF2), mRNA
NM 005368	Homo sapiens myoglobin (MB), mRNA
NM 005363	Homo sapiens melanoma antigen, family A, 6 (MAGEA6), mRNA
NM 005362	Homo sapiens melanoma antigen, family A, 3 (MAGEA3), mRNA
NM 005361	Homo sapiens melanoma antigen, family A, 2 (MAGEA2), mRNA
NM 005475	Homo sapiens lymphocyte adaptor protein (LNK), mRNA
	Homo sapiens lipase, hormone-sensitive (LIPE), mRNA
NM_005357	Homo sapiens lymphocyte-specific protein tyrosine kinase (LCK), mRNA
NM_005356	Homo sapiens potassium voltage-gated channel, Isk-related family, member 3
NM_005472	(KCNE3), mRNA
NTM 005405	Homo sapiens solute carrier family 17 (sodium phosphate), member 4
NM_005495	(SLC17A4), mRNA
NTM 005456	Homo sapiens mitogen-activated protein kinase 8 interacting protein 1
NM_005456	(MAPK8IP1), mRNA
NM_005343	Homo sapiens v-Ha-ras Harvey rat sarcoma viral oncogene homolog (HRAS),
NWI_005343	mRNA
NM_005342	Homo sapiens high-mobility group (nonhistone chromosomal) protein 4
NM_003342	(HMG4), mRNA
NM 005341	Homo sapiens GLI-Kruppel family member HKR3 (HKR3), mRNA
	Homo sapiens dell'-Krupper taminy member 11120 (1122); 11
NM_005337_	Homo sapiens hyperpolarization activated cyclic nucleotide-gated potassium
NM_005477	channel 4 (HCN4), mRNA
NM 005335	Homo sapiens hematopoietic cell-specific Lyn substrate 1 (HCLS1), mRNA
NM 005334	Homo sapiens host cell factor C1 (VP16-accessory protein) (HCFC1), mRNA
	Homo sapiens holocytochrome c synthase (cytochrome c heme-lyase) (HCCS),
NM_005333	mRNA
NM 005328	Homo sapiens hyaluronan synthase 2 (HAS2), mRNA
	Homo sapiens L-3-hydroxyacyl-Coenzyme A dehydrogenase, short chain
NM_005327	(HADHSC), mRNA
NM 005324	Homo sapiens H3 histone, family 3B (H3.3B) (H3F3B), mRNA
	Homo sapiens H1 histone family, member 4 (H1F4), mRNA
NM_005321	Homo sapiens H1 histone family, member 3 (H1F3), mRNA
NM_005320	
NM_005319	Homo sapiens H1 histone family, member 2 (H1F2), mRNA
NM_005325	Homo sapiens H1 histone family, member 1 (H1F1), mRNA
NM_005318	Homo sapiens H1 histone family, member 0 (H1F0), mRNA
NM_005459	Homo sapiens guanylate cyclase activator 1C (GUCA1C), mRNA
NM_005316	Homo sapiens general transcription factor IIH, polypeptide 1 (62kD subunit)
	(GTF2H1), mRNA
NM_005315	Homo sapiens goosecoid-like (GSCL), mRNA
NM_005314	Homo sapiens gastrin-releasing peptide receptor (GRPR), mRNA
NM_005313	Homo sapiens glucose regulated protein, 58kD (GRP58), mRNA
NM_005312	Homo sapiens guanine nucleotide-releasing factor 2 (specific for crk proto-
	oncogene) (GRF2), mRNA
NM_005311	Homo sapiens growth factor receptor-bound protein 10 (GRB10), mRNA
NM_005309	Homo sapiens glutamic-pyruvate transaminase (alanine aminotransferase)
!	(GPT), mRNA

	Linear 5 (GPPK 5) mRNA
VM 005308	Homo sapiens G protein-coupled receptor kinase 5 (GPRK5), mRNA
	Trans Carotein-counted receptor o (Or No), mix 12
	C protein coupled receptor / (Urix/), IIII(1)
NM 005284	C protein counled receptor 0 (OI NO), IIIX 121
NM 005458	Comptoin counted receptor of (ULR) 1, IIII 121
NM 005282	C protein counted receptor 4 (Or N4), mour
NM 005306	TY Consider Counted receptor 45 (Of R45), find 17
NM 005305	vi
NM 005304	G protein-coupled receptor 41 (UFN41), IIII(171)
NM 005303	VI remiens G protein-coupled receptor 40 (UFR40), IIII VI
NM_005281	Homo sapiens G protein-coupled receptor 3 (endothelin receptor type B-like)  Homo sapiens G protein-coupled receptor 37 (endothelin receptor type B-like)
NM_005302	(CDD 27) DXIA
27.6.005201	ry Greatein-coupled recentor 35 (GPR35), IIIRNA
NM_005301	Try Constein-counted receptor 34 (UI N34), illicity
NM_005300	G protein counled recentor 31 (UFR31), mid 1/2
NM_005299	Homo sapiens G protein-coupled receptor 25 (GPR25), mRNA  Homo sapiens G protein-coupled receptor 25 (GPR24), mRNA
NM_005298	Homo sapiens G protein-coupled receptor 24 (GPR24), mRNA  Homo sapiens G protein-coupled receptor 24 (GPR24), mRNA
NM_005297	Homo sapiens G protein-coupled receptor 23 (GPR23), mRNA  Homo sapiens G protein-coupled receptor 23 (GPR23), mRNA
NM_005296	Homo sapiens G protein-coupled receptor 22 (GPR22), mRNA  Homo sapiens G protein-coupled receptor 22 (GPR21), mRNA
NM_005295	Homo sapiens G protein-coupled receptor 22 (GPR21), mRNA
NM_005294	Homo sapiens G protein-coupled receptor 21 (GPR21), mRNA  Homo sapiens G protein-coupled receptor 20 (GPR20), mRNA
NM 005293	Homo sapiens G protein-coupled receptor 20 (GPR20), mRNA  Homo sapiens G protein-coupled receptor 21 (GPR1), mRNA
NM 005279	Homo sapiens G protein-coupled receptor 1 (GPR1), mRNA  Homo sapiens G protein-coupled receptor 1 (GPR17), mRNA
NM 005291	Homo sapiens G protein-coupled receptor 17 (GPR17), mRNA  Homo sapiens G protein-coupled receptor 15 (GPR15), mRNA
NM 005290	TI comions G protein-counted receptor 15 (ULK15), III. 1
NM 005288	C 4
NM 005276	· 1 phochhate den votovellase I (soluble) (SI 2 2);
NM 005275	
NM_005274	Homo sapiens guanine nucleotide binding protein (G protein), gamma
NM_005273	Homo sapiens guanine nucleotide binding protein (G protein), beta polypepines 2
	(GNB2), mRNA  Homo sapiens glutamate dehydrogenase 1 (GLUD1), mRNA  (GLUD1), mRNA
NM_005271	Homo sapiens glutamate denydrogenase i (GHODI), included Homo sapiens glioma-associated oncogene homolog (zinc finger protein) (GLI),
NM_005269	
NM_005264	Warre copiens GDNF family receptor alpha 1 (GFRA1), mRNA
NM 005263	Homo caniens growth factor independent 1 (0111), mid 12
NM 005256	TT-me conjens growth arrest-specific 2 (GAS2), IIINNA
NM 005255	Home saniens cyclin G associated kinase (GAK), IIIKNA
	Home seniens FOS-like antigen 2 (FOSL2), mRNA
NM_005253	
NM_005249	
NM_005251	
37 t 005046	Toling corooms With (V-191) Unicognic numbers
NM_005248	
	- 1 - 1 - 1 - 1 - 1 - 1 - 1 - 1 - 1 - 1
NM_005246	
	(FER), mRNA  Homo sapiens nuclear receptor subfamily 2, group F, member 6 (NR2F6),
NM_005234	Homo sapiens nuclear receptor suctamely =, 8-1-4
	mRNA  Homo sapiens EphA3 (EPHA3), mRNA
NM_00523	t and the same of
NM_00523	1 Homo sapiens ems1 sequence (maintain) tumor associated (p80/85 src substrate) (EMS1), mRNA
	associated (p80/83 Sic substrate) (Eliter); Alectic

NM_005227	Homo sapiens ephrin-A4 (EFNA4), mRNA
NM_005223	Homo sapiens deoxyribonuclease I (DNASE1), mRNA
NM_005222	Homo sapiens distal-less homeo box 6 (DLX6), mRNA
NM_005220	Homo sapiens distal-less homeo box 3 (DLX3), mRNA
NM_005216	Homo sapiens dolichyl-diphosphooligosaccharide-protein glycosyltransferase
. —	(DDOST), mRNA
NM 005215	Homo sapiens deleted in colorectal carcinoma (DCC), mRNA
NM_005436	Homo sapiens DNA segment, single copy, probe pH4 (transforming sequence,
	thyroid-1, (D10S170), mRNA
NM_005214	Homo sapiens cytotoxic T-lymphocyte-associated protein 4 (CTLA4), mRNA
NM_005213	Homo sapiens cystatin A (stefin A) (CSTA), mRNA
NM_005492	Homo sapiens cystatin 8 (cystatin-related epididymal specific) (CST8), mRNA
NM_005212	Homo sapiens casein, kappa (CSN10), mRNA
NM 005211	Homo sapiens colony stimulating factor 1 receptor, formerly McDonough feline
_	sarcoma viral (v-fms) oncogene homolog (CSF1R), mRNA
NM 005204	Homo sapiens mitogen-activated protein kinase kinase kinase 8 (MAP3K8),
_	mRNA
NM 005200	Homo sapiens cell matrix adhesion regulator (CMAR), mRNA
NM 005195	Homo sapiens CCAAT/enhancer binding protein (C/EBP), delta (CEBPD),
_	mRNA
NM 005194	Homo sapiens CCAAT/enhancer binding protein (C/EBP), beta (CEBPB),
_	mRNA
NM 005193	Homo sapiens caudal type homeo box transcription factor 4 (CDX4), mRNA
NM 005191	Homo sapiens CD80 antigen (CD28 antigen ligand 1, B7-1 antigen) (CD80),
_	mRNA
NM 005188	Homo sapiens Cas-Br-M (murine) ecotropic retroviral transforming sequence
-	(CBL), mRNA
NM_005185	Homo sapiens calmodulin-like 3 (CALML3), mRNA
NM_005184	Homo sapiens calmodulin 3 (phosphorylase kinase, delta) (CALM3), mRNA
NM_005483	Homo sapiens chromatin assembly factor 1, subunit A (p150) (CHAF1A),
	mRNA
NM_005441	Homo sapiens chromatin assembly factor 1, subunit B (p60) (CHAF1B), mRNA
NM_005183	Homo sapiens calcium channel, voltage-dependent, alpha 1F subunit
	(CACNA1F), mRNA
NM_005182	Homo sapiens carbonic anhydrase VII (CA7), mRNA
NM_005448	Homo sapiens bone morphogenetic protein 15 (BMP15), mRNA
NM_005178	Homo sapiens B-cell CLL/lymphoma 3 (BCL3), mRNA
NM_005177	Homo sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump) non-
_	catalytic accessory protein 1A (110/116kD) (ATP6N1A), mRNA
NM_005174	Homo sapiens ATP synthase, H+ transporting, mitochondrial F1 complex,
	gamma polypeptide 1 (ATP5C1), mRNA
NM_005173	Homo sapiens ATPase, Ca++ transporting, ubiquitous (ATP2A3), mRNA
NM 005171	Homo sapiens activating transcription factor 1 (ATF1), mRNA
NM 005167	Homo sapiens ras homolog gene family, member C (ARHC), mRNA
NM 005166	Homo sapiens amyloid beta (A4) precursor-like protein 1 (APLP1), mRNA
NM 005165	Homo sapiens aldolase C, fructose-bisphosphate (ALDOC), mRNA
NM 005163	Homo sapiens v-akt murine thymoma viral oncogene homolog 1 (AKT1),
	mRNA
NM 005161	Homo sapiens angiotensin receptor-like 1 (AGTRL1), mRNA
NM 005095	Homo sapiens zinc finger protein 262 (ZNF262), mRNA
NM 005096	Homo sapiens zinc finger protein 261 (ZNF261), mRNA
NM 005081	Homo sapiens zinc finger protein 142 (clone pHZ-49) (ZNF142), mRNA
14141 002001	1 Acome septems Zine miger protein 1-12 (closed pinz 15) (Zini 1-12), mid-11

PCT/US03/05028 WO 03/074654

WO 03/0/4054	o 10 LDhunit
	Homo sapiens thyroid hormone receptor-associated protein, 240 kDa subunit
NM_005121	
2.5050	
NM_005079	Homo sapiens tumor protein D52 (TPD52), interest.  Homo sapiens peptidoglycan recognition protein (PGLYRP), mRNA  Homo sapiens peptidoglycan recognition protein (PGLYRP), mRNA  Homo sapiens peptidoglycan recognition protein (PGLYRP), mRNA
NM_005091	Homo sapiens peptidoglycan recognition protein (1922) Homo sapiens tumor necrosis factor (ligand) superfamily, member 18
NM_005092	Homo sapiens tuntor neorests 2
	(TNFSF18), mRNA Homo sapiens tumor necrosis factor (ligand) superfamily, member 15
NM_005118	Homo sapiens tumor neorosis zarra (o
	(TNFSF15), mRNA Homo sapiens tumorous imaginal discs (Drosophila) homolog (TID1), mRNA  Homo sapiens tumorous imaginal discs (Drosophila) homolog (TID1), mRNA
NM_005147	Homo sapiens tumorous magnine discovery, mRNA  Homo sapiens contactin 2 (axonal) (CNTN2), mRNA  Homo sapiens contactin 2 (axonal) (CNTN2), mRNA
NM_005076	Homo sapiens contactin 2 (axonal) (CN1N2), filed 12  Homo sapiens solute carrier family 23 (nucleobase transporters), member 1
NM_005116	Homo sapiens solute carrier family 20 (
	(SLC23A1), mRNA Homo sapiens solute carrier family 4, anion exchanger, member 3 (SLC4A3),
NM_005070	Homo sapiens solute carrier family 13
	mRNA Homo sapiens solute carrier family 17 (sodium phosphate), member 1
NM_005074	Homo sapiens solute carrier ranny 17 (course)
	(SLC17A1), mRNA  Homo sapiens solute carrier family 15 (oligopeptide transporter), member 1
NM_005073	Homo sapiens solute carrier failing 13 (ongoth
	(SLC15A1), mRNA Homo sapiens solute carrier family 12 (potassium/chloride transporters), member
NM_005072	Homo sapiens solute carrier family 12 (poussessess)
_	4 (SLC12A4), mRNA  Homo sapiens stearoyl-CoA desaturase (delta-9-desaturase) (SCD), mRNA  Homo sapiens stearoyl-CoA desaturase (delta-9-desaturase) (SCD), mRNA
NM 005063	Homo sapiens stearoyl-CoA desaurase (dotta ) GORC), mRNA
NM 005060	Transa conjens RAR-related orbital receptor 5 (2-2-2)
NM 005059	Homo sapiens relaxin 2 (H2) (RLN2), history
NM_005045	Homo sapiens reelin (RELN), mRNA  Homo sapiens RNA binding motif protein, Y chromosome, family 1, member A1
NM_005058	Homo sapiens RNA binding motif protein, 1 chi shada
14141_00000	(RBMY1A1), mRNA  Homo sapiens ras-related C3 botulinum toxin substrate 3 (rho family, small GTP
NM_005052	Homo sapiens ras-related C3 botulinum toxin substrate 5 (The American
14141_005052	
NM_005051	
NM 005048	·tharroad normanne (CUUDIOL 2 (* *********************************
NM 005044	Homo sapiens paratnyloid normens yet protein kinase, X-linked (PRKX), mRNA  Homo sapiens protein kinase, X-linked (PRKX), mRNA
NM 005043	
NM 005042	
NM 00504	Homo sapiens proline-rich protein Hacht successful (PRF1), mRNA Homo sapiens perforin 1 (preforming protein) (PRF1), mRNA  Homo sapiens proline-rich protein Hacht successful (PRF1), mRNA
	Homo sapiens perforin 1 (preforming protein) (TRF1), mRNA Homo sapiens prolylcarboxypeptidase (angiotensinase C) (PRCP), mRNA Homo sapiens prolylcarboxypeptidase (angiotensinase C) (PRCP), mRNA
NM 005040	Homo sapiens prolylcarboxypeptidase (angioteinmann) Homo sapiens proline-rich protein BstNI subfamily 1 (PRB1), mRNA (cyclophilin D) (PPID), mRNA
NM_00503	Homo sapiens proline-rich protein BstN1 subtainity 1 (1832), mRNA Homo sapiens peptidylprolyl isomerase D (cyclophilin D) (PPID), mRNA Homo sapiens peptidylprolyl isomerase D (cyclophilin D) (PPID), mRNA
NM_00503	Homo sapiens peptidylprolyl isomerase D (cyclopinini D) (122), mRNA Homo sapiens paired-like homeodomain transcription factor 3 (PITX3), mRNA like is in the same of the same
NM_00502	
NM_00502	beta) (PIK3R2), mRNA
- T C 00500	1 1 am ocitive 4 - Killiase Caldivity of the F
NM_00502	mDNA (CNDD2)
	mRNA Homo sapiens ectonucleotide pyrophosphatase/phosphodiesterase 3 (ENPP3),
NM_00502	Holito sapiono cotomaso a la mana a m
	mRNA  Homo sapiens phosphodiesterase 1A, calmodulin-dependent (PDE1A), mRNA  local death 1 (PDCD1), mRNA
NM_0050	Homo sapiens prospriodiesterase 11; established Homo sapiens programmed cell death 1 (PDCD1), mRNA  Homo sapiens programmed cell death 1 (PDCD1), mRNA
NM_0050	
NM_0050	1 mm misma muslaanama /14k1/10/10/10/10/10/10/10/10/10/10/10/10/10
NM_0050	
NM_0051	Homo sapiens nucleobindin 2 (NUCB2), mRNA  Homo sapiens nucleobindin 2 (NUCB2), mRNA  Homo sapiens nucleobindin 2 (NUCB2), mRNA
NM_0050	
NM_0050	
NM 0050	11 Homo sapiens nuclear respiratory factor

1: AD(D() D) IA		
NM 005007 Homo sapiens non-metastatic cells 4, protein expressed in (NMEA), mRNA NM 005007 Homo sapiens nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor-like 1 (NFKBIL1), mRNA NM 005004 Homo sapiens NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 8 (19kD, ASHI) (NDUFB8), mRNA NM 005001 Homo sapiens NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 7 (14.5kD, B14.5a) (NDUFA7), mRNA NM 005007 Homo sapiens leucine-rich, glioma inactivated 1 (LGI1), mRNA NM 005007 Homo sapiens leucine-rich, glioma inactivated 1 (LGI1), mRNA NM 004983 Homo sapiens leucine-rich, glioma inactivated 1 (LGI1), mRNA NM 004983 Homo sapiens leucine-rich, glioma inactivated 1 (LGI1), mRNA NM 004983 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 9 (KCND9), mRNA NM 004982 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 9 (KCND9), mRNA NM 005090 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 5 (KCNJ8), mRNA NM 005136 Homo sapiens potassium vinwardly-rectifying channel, subfamily J, member 4 (KCND4), mRNA NM 005136 Homo sapiens potassium voltage-gated channel, shal-related family, member 2 (KCND2), mRNA NM 004980 Homo sapiens potassium voltage-gated channel, shal-related subfamily, member 3 (KCND3), mRNA NM 004970 Homo sapiens potassium voltage-gated channel, Shal-related subfamily, member 4 (KCNCA), mRNA NM 004970 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCND1), mRNA Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCND1), mRNA Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCND1), mRNA Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCND1), mRNA Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCND1), mRNA Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCND1), mRNA Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1	NM 005010	Homo sapiens neuronal cell adhesion molecule (NRCAM), mRNA
NM_005007  Homo sapiens nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor-like 1 (NFKEBL1), mRNA  NM_005004  Homo sapiens NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 8 (19kD, ASHI) (NDUFB8), mRNA  NM_005001  Homo sapiens MADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 7 (14.5kD, B14.5a) (NDUFA7), mRNA  NM_004988  MM_004988  Homo sapiens melanoma antigen, family A, 1 (directs expression of antigen MZ2-E) (MAGEA1), mRNA  NM_004981  Homo sapiens leucine-rich, glioma inactivated 1 (LGI1), mRNA  NM_004983  Homo sapiens sheisin family member 5A (KEFSA), mRNA  NM_004984  Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 9 (KCNI9), mRNA  NM_004983  Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 9 (KCNI5), mRNA  NM_004981  Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 5 (KCNI5), mRNA  NM_004981  Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 5 (KCNI5), mRNA  NM_004981  Homo sapiens potassium voltage-gated channel, Isk-related family, member 2 (KCNE2), mRNA  NM_004978  Homo sapiens potassium voltage-gated channel, Shal-related subfamily, member 3 (KCND1), mRNA  NM_004978  Homo sapiens potassium voltage-gated channel, Shal-related subfamily, member 1 (KCNO1), mRNA  NM_004978  Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 3 (KCNC3), mRNA  NM_004978  Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNC1), mRNA  NM_004978  Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNB1), mRNA  NM_004979  Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNB1), mRNA  NM_004970  Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNB1), mRNA  NM_004970  Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNB1), mRNA  Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNB1), mRNA  NM_004970  Ho	NM 005009	Homo sapiens non-metastatic cells 4, protein expressed in (NME4), mRNA
inhibitor-like 1 (NFKBIL1), mRNA NM_005004 Homo sapiens NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 8 (19kD, ASHI) (NDUFB8), mRNA NM_005001 Homo sapiens MADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 7 (14.5kD, B14.5a) (NDUFA7), mRNA NM_004988 Homo sapiens melanoma antigen, family A, 1 (directs expression of antigen M22-E) (MAGEA1), mRNA NM_004981 Homo sapiens leucine-rich, glioma inactivated 1 (LGII), mRNA NM_004983 Homo sapiens leucine-rich, glioma inactivated 1 (LGII), mRNA NM_004983 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 9 (KCNJ9), mRNA NM_004983 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 9 (KCNJ9), mRNA NM_004981 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 8 (KCNJ8), mRNA NM_004982 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 5 (KCNJ8), mRNA NM_0004981 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 5 (KCNJ4), mRNA NM_004981 Homo sapiens potassium voltage-gated channel, Isk-related family, member 2 (KCNE2), mRNA NM_004970 Homo sapiens potassium voltage-gated channel, Shal-related subfamily, member 3 (KCND3), mRNA NM_004979 Homo sapiens potassium voltage-gated channel, Shal-related family, member 1 (KCND1), mRNA NM_004978 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 4 (KCND4), mRNA NM_004977 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNC1), mRNA NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCND1), mRNA NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCND1), mRNA NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNB1), mRNA NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNB1), mRNA NM_004976 Homo sapiens spotassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNB1), mRNA NM_004976 Homo sapiens spotassium vol	NM 005007	Homo sapiens nuclear factor of kappa light polypeptide gene enhancer in B-cells
NM_005004 Homo sapiens NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 8 (19kD, ASHI) (NDUFB8), mRNA  NM_005001 Homo sapiens NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 7 (14.5kD, B14.5a) (NDUFA7), mRNA  NM_004988 Homo sapiens melanoma antigen, family A, 1 (directs expression of antigen MZ2-E) (MAGEA1), mRNA  NM_004981 Homo sapiens leucine-rich, glioma inactivated 1 (LGI1), mRNA  NM_004984 Homo sapiens leucine-rich, glioma inactivated 1 (LGI1), mRNA  NM_004984 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 9 (KCNJ9), mRNA  NM_004984 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 8 (KCNJS), mRNA  NM_004982 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 5 (KCNJ5), mRNA  NM_004981 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 5 (KCNJ4), mRNA  NM_005136 Homo sapiens potassium voltage-gated channel, lsk-related family, member 2 (KCNE2), mRNA  NM_004980 Homo sapiens potassium voltage-gated channel, Shal-related subfamily, member 3 (KCND3), mRNA  NM_004978 Homo sapiens potassium voltage-gated channel, Shal-related family, member 1 (KCND1), mRNA  NM_004978 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 4 (KCNC4), mRNA  NM_004978 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNB1), mRNA  NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNB1), mRNA  NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNB1), mRNA  NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNB1), mRNA  NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNB1), mRNA  NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNB1), mRNA  NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNB1), mRNA  NM_004976 Homo sapiens potassium voltage-gated	_	inhibitor-like 1 (NFKBIL1), mRNA
ASHI) (NDUFB8), mRNA  NM_004988 Homo sapiens NADH dehydrogenase (ubiquinone) 1 alipha subcomplex, 7 (14.5kD, B14.5a) (NDUFA7), mRNA  NM_004988 Homo sapiens belanoma antigen, family A, 1 (directs expression of antigen MZ2-E) (MAGEA1), mRNA  NM_004981 Homo sapiens leucine-rich, glioma inactivated 1 (LGII), mRNA  NM_004983 Homo sapiens kinesin family member 5A (KIF5A), mRNA  NM_004984 Homo sapiens sotassium inwardly-rectifying channel, subfamily J, member 9 (KCNJ9), mRNA  NM_004982 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 8 (KCNJ8), mRNA  NM_004982 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 5 (KCNJ5), mRNA  NM_004981 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 4 (KCNJ4), mRNA  NM_004981 Homo sapiens potassium voltage-gated channel, sk-related family, member 2 (KCNE2), mRNA  NM_004980 Homo sapiens potassium voltage-gated channel, Shal-related subfamily, member 3 (KCND3), mRNA  NM_004970 Homo sapiens potassium voltage-gated channel, Shal-related family, member 1 (KCND1), mRNA  NM_004977 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 3 (KCNG1), mRNA  NM_004977 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNC1), mRNA  NM_004977 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNC1), mRNA  NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNE1), mRNA  NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNE1), mRNA  NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNE1), mRNA  NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNE1), mRNA  NM_004969 Homo sapiens spotassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNE1), mRNA  NM_004960 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNE1), mRNA  NM_004960 Homo sapiens spotassium	NM 005004	Homo sapiens NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 8 (19kD,
NM_004981 Homo sapiens NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 7 (14.5kD, B14.5a) (NDUFA7), mRNA NM_004988 Homo sapiens melanoma antigen, family A, 1 (directs expression of antigen MZ2-E) (MAGEA1), mRNA NM_005097 Homo sapiens leucine-rich, glioma inactivated 1 (LGII), mRNA NM_004984 Homo sapiens leucine-rich, glioma inactivated 1 (LGII), mRNA NM_004984 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 9 (KCNI9), mRNA NM_004982 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 8 (KCNI8), mRNA NM_004980 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 5 (KCNI5), mRNA NM_004981 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 4 (KCNI4), mRNA NM_005136 Homo sapiens potassium voltage-gated channel, sk-related family, member 2 (KCNE2), mRNA NM_004980 Homo sapiens potassium voltage-gated channel, Shal-related subfamily, member 3 (KCND3), mRNA NM_004979 Homo sapiens potassium voltage-gated channel, Shal-related family, member 1 (KCND1), mRNA NM_004978 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 3 (KCNO3), mRNA NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNCI), mRNA NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNC1), mRNA NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNS1), mRNA NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNS1), mRNA NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNS1), mRNA NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNS1), mRNA NM_004960 Homo sapiens haptoglobin (HP), mRNA NM_004961 Homo sapiens spatial maligner potage potage channel, Shaw-related subfamily, member 1 (KCNS1), mRNA NM_004961 Homo sapiens papting-bindigner potage potage channel, Shaw-related subfamily, member 1 (KCNS1), mRNA NM_00510	" <del>-</del>	ASHI) (NDUFB8), mRNA
(14.5kD, B14.5a) (NDUFA7), mRNA  NM_004988 Homo sapiens melanoma antigen, family A, 1 (directs expression of antigen MZ2-E) (MAGEA1), mRNA  NM_003097 Homo sapiens leucine-rich, glioma inactivated 1 (LGI1), mRNA  NM_004984 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 9 (KCNJ9), mRNA  NM_004982 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 8 (KCNJ9), mRNA  NM_004982 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 8 (KCNJ8), mRNA  NM_004980 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 5 (KCNJ5), mRNA  NM_004981 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 5 (KCNJ4), mRNA  NM_004981 Homo sapiens potassium voltage-gated channel, subfamily J, member 2 (KCNE2), mRNA  NM_004980 Homo sapiens potassium voltage-gated channel, Shal-related family, member 2 (KCND3), mRNA  NM_004980 Homo sapiens potassium voltage-gated channel, Shal-related subfamily, member 3 (KCND3), mRNA  NM_004970 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 4 (KCNC4), mRNA  NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 3 (KCNC3), mRNA  NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNE1), mRNA  NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNE1), mRNA  NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNE1), mRNA  NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNE1), mRNA  NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNE1), mRNA  NM_004969 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNE1), mRNA  NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNE1), mRNA  NM_004969 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNE1), mRNA  N	NM 005001	Homo sapiens NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 7
NM_004988	1111_00000	(14.5kD, B14.5a) (NDUFA7), mRNA
MZ2-E) (MAGEA1), mRNA NM 004984 Homo sapiens leucine-rich, glioma inactivated 1 (LGII), mRNA NM 004984 Homo sapiens leucine-rich, glioma inactivated 1 (LGII), mRNA NM_004984 Homo sapiens kinesin family member 5A (KIF5A), mRNA NM_004982 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 9 (KCNJ9), mRNA NM_004982 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 8 (KCNJ8), mRNA NM_004981 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 5 (KCNJ5), mRNA NM_004981 Homo sapiens potassium voltage-gated channel, subfamily J, member 4 (KCNJ4), mRNA NM_005136 Homo sapiens potassium voltage-gated channel, Isk-related family, member 2 (KCNE2), mRNA NM_004980 Homo sapiens potassium voltage-gated channel, Shal-related subfamily, member 3 (KCND3), mRNA NM_004979 Homo sapiens potassium voltage-gated channel, Shal-related family, member 1 (KCND1), mRNA NM_004978 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 3 (KCNC3), mRNA NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 3 (KCNC3), mRNA NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNC1), mRNA NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNC1), mRNA NM_004961 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNB1), mRNA NM_004965 Homo sapiens haptoglobin (HP), mRNA NM_004966 Homo sapiens haptoglobin (HP), mRNA NM_004966 Homo sapiens haptoglobin (HP), mRNA NM_005130 Homo sapiens heparin-binding growth factor binding protein (HBP17), mRNA NM_005141 Homo sapiens guanylate cyclase 2C (heat stable enterotoxin receptor) (GUCY2C), mRNA NM_005100 Homo sapiens guanylate cyclase 2C (heat stable enterotoxin receptor) (GUCY2C), mRNA NM_005101 Homo sapiens guanylate cyclase 2C (heat stable enterotoxin receptor) (GUCY2C), mRNA NM_005104 Homo sapiens guanylate cyclase 2C (beat stable enterotoxin receptor) (GUCY2C), mRNA NM_005104 Homo sapie	NM 004988	Homo sapiens melanoma antigen, family A, 1 (directs expression of antigen
Mono Sapiens leucine-rich, glioma inactivated 1 (LGII), mRNA	1111_00 1500	MZ2-E) (MAGEA1), mRNA
NM         004984         Homo sapiens kinesin family member 5A (KIFSA), mRNA           NM_004983         Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 9 (KCNJS), mRNA           NM_004982         Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 8 (KCNJS), mRNA           NM_000890         Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 5 (KCNJS), mRNA           NM_004981         Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 4 (KCNJA), mRNA           NM_005136         Homo sapiens potassium voltage-gated channel, Isk-related family, member 2 (KCNE2), mRNA           NM_004980         Homo sapiens potassium voltage-gated channel, Shal-related subfamily, member 3 (KCND3), mRNA           NM_004970         Homo sapiens potassium voltage-gated channel, Shak-related family, member 1 (KCND1), mRNA           NM_004978         Homo sapiens potassium voltage-gated channel, Shak-related subfamily, member 3 (KCNC3), mRNA           NM_004976         Homo sapiens potassium voltage-gated channel, Shak-related subfamily, member 1 (KCNE1), mRNA           NM_004975         Homo sapiens potassium voltage-gated channel, Shak-related subfamily, member 1 (KCNE1), mRNA           NM_004969         Homo sapiens insulin-degrading enzyme (IDE), mRNA           NM_004969         Homo sapiens sinsulin-degrading enzyme (IDE), mRNA           NM_005103         Homo sapiens heparin-binding growth factor binding protein (HBP17), mRNA	NM 005097	Homo saniens leucine-rich, glioma inactivated 1 (LGII), mRNA
NM_004982 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 9 (KCNJ9), mRNA NM_004982 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 8 (KCNJ8), mRNA NM_00890 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 5 (KCNJ5), mRNA NM_004981 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 4 (KCNJ4), mRNA NM_005136 Homo sapiens potassium voltage-gated channel, Isk-related family, member 2 (KCNE2), mRNA NM_004980 Homo sapiens potassium voltage-gated channel, Shal-related subfamily, member 3 (KCND3), mRNA NM_004979 Homo sapiens potassium voltage-gated channel, Shal-related family, member 1 (KCND1), mRNA NM_004978 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 4 (KCNC4), mRNA NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 3 (KCNC3), mRNA NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNB1), mRNA NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNB1), mRNA NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNB1), mRNA NM_004969 Homo sapiens insulin-degrading enzyme (IDE), mRNA NM_004961 Homo sapiens insulin-degrading enzyme (IDE), mRNA NM_004961 Homo sapiens haptoglobin (HP), mRNA NM_004963 Homo sapiens haptoglobin (HP), mRNA NM_005100 Homo sapiens haptoglobin (PR), mRNA NM_005101 Homo sapiens potassium voltage-gated stable enterotoxin receptor (GUCY2C), mRNA NM_005101 Homo sapiens gataric intrinsic factor (vitamin B synthesis) (GIF), mRNA NM_005104 Homo sapiens gataric intrinsic factor (vitamin B synthesis) (GIF), mRNA NM_005104 Homo sapiens gastric intrinsic factor (vitamin B synthesis) (GIF), mRNA NM_005104 Homo sapiens gastric intrinsic factor (vitamin B synthesis) (GIF), mRNA NM_005104 Homo sapiens gustric intrinsic factor (vitamin B synthesis) (GIF), mRNA NM_005104 Homo sapiens gustric intrinsic factor (vitamin B synthesis) (GIF), mRN		Homo sapiens kinesin family member 5A (KIF5A), mRNA
K(KCNJ9), mRNA   Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 8 (KCNJS), mRNA   Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 5 (KCNJS), mRNA   Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 4 (KCNJ4), mRNA   Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 4 (KCNJ4), mRNA   Homo sapiens potassium voltage-gated channel, Isk-related family, member 2 (KCNE2), mRNA   Homo sapiens potassium voltage-gated channel, Shal-related subfamily, member 3 (KCND3), mRNA   Homo sapiens potassium voltage-gated channel, Shal-related family, member 1 (KCND1), mRNA   Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 4 (KCNC4), mRNA   Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 3 (KCNC3), mRNA   Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 3 (KCNC1), mRNA   Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNB1), mRNA   Homo sapiens potassium voltage-gated channel, Shab-related subfamily, member 1 (KCNB1), mRNA   Homo sapiens potassium voltage-gated channel, Shab-related subfamily, member 1 (KCNB1), mRNA   Homo sapiens insulin-degrading enzyme (IDE), mRNA   Homo sapiens insulin-degrading enzyme (IDE), mRNA   Homo sapiens haptoglobin (HP), mRNA   Homo sapiens haptoglobin (HP), mRNA   Homo sapiens gaspiens haptoglobin (HP), mRNA   Homo sapiens gaspiens gaspiens growth factor binding protein (HBP17), mRNA   Homo sapiens gaspiens gaspiens golgin subfamily a, 5 (GOLGA5), mRNA   Homo sapiens gaspiens galatoantigen, golgin subfamily a, 5 (GOLGA5), mRNA   Homo sapiens gastric intrinsic factor (vitamin B synthesis) (GIF), mRNA   Homo sapiens gastric intrinsic factor (vitamin B synthesis) (GIF), mRNA   Homo sapiens gustrien intrinsic factor (vitamin B synthesis) (GIF), mRNA   Homo sapiens gustrien intrinsic factor (vitamin B synthesis) (HPT), mRNA   Homo sapiens gustrien intrinsic factor (vitamin B synthesis) (HPT), mRN		Homo sapiens notassium inwardly-rectifying channel, subfamily J, member 9
NM_004982 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 8 (KCNI8), mRNA  NM_004981 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 5 (KCNJ5), mRNA  NM_004981 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 4 (KCNJ4), mRNA  NM_005136 Homo sapiens potassium voltage-gated channel, Isk-related family, member 2 (KCNE2), mRNA  NM_004980 Homo sapiens potassium voltage-gated channel, Shal-related subfamily, member 3 (KCND3), mRNA  NM_004979 Homo sapiens potassium voltage-gated channel, Shal-related family, member 1 (KCND1), mRNA  NM_004978 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 4 (KCNC4), mRNA  NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 3 (KCNC3), mRNA  NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNC1), mRNA  NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNB1), mRNA  NM_004969 Homo sapiens potassium voltage-gated channel, Shab-related subfamily, member 1 (KCNB1), mRNA  NM_004969 Homo sapiens haptoglobin (HP), mRNA  NM_005143 Homo sapiens haptoglobin (HP), mRNA  NM_004965 Homo sapiens haptoglobin (HP), mRNA  NM_005100 Homo sapiens heparin-binding growth factor binding protein (HBP17), mRNA  NM_005100 Homo sapiens keparin-binding growth factor binding protein (HBP17), mRNA  NM_005101 Homo sapiens gaunylate cyclase 2C (heat stable enterotoxin receptor) (GUCY2C), mRNA  NM_005101 Homo sapiens guanylate cyclase 2C (heat stable enterotoxin receptor) (GUCY2C), mRNA  NM_005100 Homo sapiens guantine nucleotide binding protein (Gravin) 12 (AKAP12), mRNA  NM_005101 Homo sapiens guanine nucleotide binding protein (Grotein), gamma 7  (GNG7), mRNA  NM_005104 Homo sapiens guanine nucleotide binding protein (Gravin), gamma 7  (GNG7), mRNA  NM_005104 Homo sapiens guanine-fructose-6-phosphate transaminase 2 (GFPT2), mRNA  NM_004969 Homo sapiens fusion, derived from t(12;16) malignant liposarcom	NM_004963	
(KCNJ8), mRNA  NM_004981 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 5 (KCNJ5), mRNA  NM_004981 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 4 (KCNJ4), mRNA  NM_005136 Homo sapiens potassium voltage-gated channel, Isk-related family, member 2 (KCNE2), mRNA  NM_004980 Homo sapiens potassium voltage-gated channel, Shal-related subfamily, member 3 (KCND3), mRNA  NM_004979 Homo sapiens potassium voltage-gated channel, Shal-related family, member 1 (KCND1), mRNA  NM_004978 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 4 (KCNC4), mRNA  NM_004977 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 3 (KCNC3), mRNA  NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNC1), mRNA  NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNC1), mRNA  NM_004975 Homo sapiens potassium voltage-gated channel, Shab-related subfamily, member 1 (KCNB1), mRNA  NM_004965 Homo sapiens insulin-degrading enzyme (IDE), mRNA  NM_004965 Homo sapiens haptoglobin (HP), mRNA  NM_004965 Homo sapiens haptoglobin (HP), mRNA  NM_005130 Homo sapiens haptoglobin (HP), mRNA  NM_005130 Homo sapiens haptoglobin (HP), mRNA  NM_005130 Homo sapiens guanylate cyclase 2C (heat stable enterotoxin receptor) (GUCY2C), mRNA  NM_005101 Homo sapiens guanylate cyclase 2C (heat stable enterotoxin receptor) (GUCY2C), mRNA  NM_005101 Homo sapiens guanine nucleotide binding protein (Gravin) 12 (AKAP12), mRNA  NM_005104 Homo sapiens guanine nucleotide binding protein (Grotein), ganuma 7 (GNG7), mRNA  NM_005104 Homo sapiens guarnine-fructose-6-phosphate transaminase 2 (GFPT2), mRNA  NM_005104 Homo sapiens fusion, derived from t(12;16) malignant liposarcoma (FUS), mRNA  NM_004969 Homo sapiens fusion, derived from t(12;16) malignant liposarcoma (FUS), mRNA	ND 4 004092	Home senions notessium inwardly-rectifying channel subfamily I member 8
NM_004981   Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 5 (KCN15), mRNA	NM_004982	
(KCNJ5), mRNA  NM_004981 Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 4 (KCNJ4), mRNA  NM_005136 Homo sapiens potassium voltage-gated channel, Isk-related family, member 2 (KCNE2), mRNA  NM_004980 Homo sapiens potassium voltage-gated channel, Shal-related subfamily, member 3 (KCND3), mRNA  NM_004979 Homo sapiens potassium voltage-gated channel, Shal-related family, member 1 (KCND1), mRNA  NM_004978 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 4 (KCNC4), mRNA  NM_004977 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 3 (KCNC3), mRNA  NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNC1), mRNA  NM_004975 Homo sapiens potassium voltage-gated channel, Shab-related subfamily, member 1 (KCNB1), mRNA  NM_004969 Homo sapiens potassium voltage-gated channel, Shab-related subfamily, member 1 (KCNB1), mRNA  NM_005143 Homo sapiens insulin-degrading enzyme (IDE), mRNA  NM_005144 Homo sapiens haptoglobin (HP), mRNA  NM_005145 Homo sapiens high-mobility group (nonhistone chromosomal) protein 14 (HMG14), mRNA  NM_005130 Homo sapiens guanylate cyclase 2C (heat stable enterotoxin receptor) (GUCY2C), mRNA  NM_005100 Homo sapiens guanylate cyclase 2C (heat stable enterotoxin receptor) (GUCY2C), mRNA  NM_005101 Homo sapiens guanine nucleotide binding protein (G protein), gamma 7 (GNG7), mRNA  NM_005145 Homo sapiens guanine nucleotide binding protein (G protein), gamma 7 (GNG7), mRNA  NM_005140 Homo sapiens guanine nucleotide binding protein (G protein), gamma 7 (GNG7), mRNA  NM_005140 Homo sapiens gastric intrinsic factor (vitamin B synthesis) (GIF), mRNA  NM_005141 Homo sapiens guanine nucleotide binding protein (G protein), gamma 7 (GNG7), mRNA  NM_005145 Homo sapiens guanine nucleotide binding protein (G protein), gamma 7 (GNG7), mRNA  NM_005145 Homo sapiens gastric intrinsic factor (vitamin B synthesis) (GIF), mRNA  NM_005145 Homo sapiens pastric intrinsic factor (vitamin B synthesis) (GIF), mRNA  N	3.D.f. 000000	(KCNJO), IIIKNA
NM_004981	MW_000890	
(KCNJ4), mRNA	27 ( 00 1001	(KCNJ5), mkNA
NM_004980	NM_004981	
(KCNE2), mRNA   Homo sapiens potassium voltage-gated channel, Shal-related subfamily, member 3 (KCND3), mRNA   Homo sapiens potassium voltage-gated channel, Shal-related family, member 1 (KCND1), mRNA   Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 4 (KCNC4), mRNA   Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 3 (KCNC3), mRNA   Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 3 (KCNC3), mRNA   Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNC1), mRNA   Homo sapiens potassium voltage-gated channel, Shab-related subfamily, member 1 (KCNB1), mRNA   Homo sapiens insulin-degrading enzyme (IDE), mRNA   MM_004969   Homo sapiens insulin-degrading enzyme (IDE), mRNA   Homo sapiens haptoglobin (HP), mRNA   MM_004965   Homo sapiens haptoglobin (HP), mRNA   MM_004965   Homo sapiens heparin-binding growth factor binding protein (HBP17), mRNA   MM_005130   Homo sapiens potassium voltage-gated channel, Shab-related subfamily   Homo sapiens guanylate cyclase 2C (heat stable enterotoxin receptor) (GUCY2C), mRNA   Homo sapiens guanylate cyclase 2C (heat stable enterotoxin receptor) (GUCY2C), mRNA   Homo sapiens golgi autoantigen, golgin subfamily a, 5 (GOLGA5), mRNA   MM_005113   Homo sapiens guanine nucleotide binding protein (G protein), gamma 7 (GNG7), mRNA   Homo sapiens guanine fuctore-6-phosphate transaminase 2 (GFPT2), mRNA   NM_00510   Homo sapiens fusion, derived from t(12;16) malignant liposarcoma (FUS), mRNA   Homo sapiens nuclear receptor subfamily 5, group A, member 1 (NR5A1), mRNA   Homo sapiens nuclear receptor subfamily 5, group A, member 1 (NR5A1), mRNA   Homo sapiens nuclear receptor subfamily 5, group A, member 1 (NR5A1), mRNA   Homo sapiens nuclear receptor subfamily 5, group A, member 1 (NR5A1), mRNA   Homo sapiens nuclear receptor subfamily 5, group A, member 1 (NR5A1), mRNA   Homo sapiens nuclear receptor subfamily 5, group A, member 1 (NR5A1), mRNA   Homo sapiens nuclear receptor		(KCNJ4), mKNA
NM_004979 Homo sapiens potassium voltage-gated channel, Shal-related subfamily, member 3 (KCND3), mRNA  NM_004979 Homo sapiens potassium voltage-gated channel, Shal-related family, member 1 (KCND1), mRNA  NM_004978 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 4 (KCNC4), mRNA  NM_004977 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 3 (KCNC3), mRNA  NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNC1), mRNA  NM_004975 Homo sapiens potassium voltage-gated channel, Shab-related subfamily, member 1 (KCNB1), mRNA  NM_004969 Homo sapiens potassium voltage-gated channel, Shab-related subfamily, member 1 (KCNB1), mRNA  NM_005143 Homo sapiens insulin-degrading enzyme (IDE), mRNA  NM_005144 Homo sapiens haptoglobin (HP), mRNA  NM_005130 Homo sapiens high-mobility group (nonhistone chromosomal) protein 14 (HMG14), mRNA  NM_004963 Homo sapiens potassium voltage-gated channel, Shab-related subfamily, member 1 (KCNB1), mRNA  NM_005100 Homo sapiens haptoglobin (HP), mRNA  NM_005100 Homo sapiens guanylate cyclase 2C (heat stable enterotoxin receptor) (GUCY2C), mRNA  NM_005113 Homo sapiens A kinase (PRKA) anchor protein (gravin) 12 (AKAP12), mRNA  NM_005145 Homo sapiens golgi autoantigen, golgin subfamily a, 5 (GOLGA5), mRNA  NM_005145 Homo sapiens gastric intrinsic factor (vitamin B synthesis) (GIF), mRNA  NM_00510 Homo sapiens gastric intrinsic factor (vitamin B synthesis) (GIF), mRNA  NM_00510 Homo sapiens fusion, derived from t(12;16) malignant liposarcoma (FUS), mRNA  NM_004960 Homo sapiens nuclear receptor subfamily 5, group A, member 1 (NR5A1), mRNA	NM_005136	
NM_004979 Homo sapiens potassium voltage-gated channel, Shal-related family, member 1 (KCND1), mRNA  NM_004978 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 4 (KCNC4), mRNA  NM_004977 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 3 (KCNC3), mRNA  NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNC1), mRNA  NM_004975 Homo sapiens potassium voltage-gated channel, Shab-related subfamily, member 1 (KCNB1), mRNA  NM_004969 Homo sapiens insulin-degrading enzyme (IDE), mRNA  NM_005143 Homo sapiens haptoglobin (HP), mRNA  NM_004965 Homo sapiens high-mobility group (nonhistone chromosomal) protein 14 (HMG14), mRNA  NM_005130 Homo sapiens heparin-binding growth factor binding protein (HBP17), mRNA  NM_005100 Homo sapiens guanylate cyclase 2C (heat stable enterotoxin receptor) (GUCY2C), mRNA  NM_005101 Homo sapiens golgi autoantigen, golgin subfamily a, 5 (GOLGA5), mRNA  NM_005145 Homo sapiens guanine nucleotide binding protein (G protein), gamma 7 (GNG7), mRNA  NM_005142 Homo sapiens gastric intrinsic factor (vitamin B synthesis) (GIF), mRNA  NM_00510 Homo sapiens gastric intrinsic factor (vitamin B synthesis) (GIF), mRNA  NM_00510 Homo sapiens gastric intrinsic factor (vitamin B synthesis) (GIF), mRNA  NM_005142 Homo sapiens gastric intrinsic factor (vitamin B synthesis) (GIF), mRNA  NM_00510 Homo sapiens gastric intrinsic factor (vitamin B synthesis) (GIF), mRNA  NM_00510 Homo sapiens fusion, derived from t(12;16) malignant liposarcoma (FUS), mRNA  NM_004959 Homo sapiens nuclear receptor subfamily 5, group A, member 1 (NR5A1), mRNA		(KCNE2), mRNA
NM_004979   Homo sapiens potassium voltage-gated channel, Shal-related family, member 1 (KCND1), mRNA   Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 4 (KCNC4), mRNA   Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 3 (KCNC3), mRNA   Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNC1), mRNA   Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNB1), mRNA   Homo sapiens potassium voltage-gated channel, Shab-related subfamily, member 1 (KCNB1), mRNA   Homo sapiens potassium voltage-gated channel, Shab-related subfamily, member 1 (KCNB1), mRNA   Homo sapiens insulin-degrading enzyme (IDE), mRNA   MM_005143   Homo sapiens haptoglobin (HP), mRNA   Homo sapiens guanylate cyclase 2C (heat stable enterotoxin receptor) (GUCY2C), mRNA   Homo sapiens golgi autoantigen, golgin subfamily a, 5 (GOLGA5), mRNA   Homo sapiens golgi autoantigen, golgin subfamily a, 5 (GOLGA5), mRNA   Homo sapiens guanine nucleotide binding protein (G protein), gamma 7 (GNG7), mRNA   Homo sapiens gastric intrinsic factor (vitamin B synthesis) (GIF), mRNA   NM_005110   Homo sapiens guatine-fructose-6-phosphate transaminase 2 (GFPT2), mRNA   NM_004960   Homo sapiens fusion, derived from t(12;16) malignant liposarcoma (FUS), mRNA   Homo sapiens nuclear receptor subfamily 5, group A, member 1 (NR5A1), mRNA   Homo sapiens nuclear receptor subfamily 5, group A, member 1 (NR5A1), mRNA   Homo sapiens nuclear receptor subfamily 5, group A, member 1 (NR5A1), mRNA   Homo sapiens nuclear receptor subfamily 5, group A, member 1 (NR5A1), mRNA   Homo sapiens nuclear receptor subfamily 5, group A, member 1 (NR5A1), mRNA   Homo sapiens nuclear receptor subfamily 5, group A, member 1 (NR5A1), mRNA   Homo sapiens nuclear receptor subfamily 5, group A, me	NM_004980	
(KCND1), mRNA  Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 4 (KCNC4), mRNA  NM_004977 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 3 (KCNC3), mRNA  NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNC1), mRNA  NM_004975 Homo sapiens potassium voltage-gated channel, Shab-related subfamily, member 1 (KCNB1), mRNA  NM_004969 Homo sapiens potassium voltage-gated channel, Shab-related subfamily, member 1 (KCNB1), mRNA  NM_004969 Homo sapiens insulin-degrading enzyme (IDE), mRNA  NM_005143 Homo sapiens haptoglobin (HP), mRNA  NM_004965 Homo sapiens high-mobility group (nonhistone chromosomal) protein 14 (HMG14), mRNA  NM_005130 Homo sapiens peparin-binding growth factor binding protein (HBP17), mRNA  NM_004963 Homo sapiens guanylate cyclase 2C (heat stable enterotoxin receptor) (GUCY2C), mRNA  NM_005100 Homo sapiens A kinase (PRKA) anchor protein (gravin) 12 (AKAP12), mRNA  NM_005113 Homo sapiens guanine nucleotide binding protein (G protein), gamma 7 (GNG7), mRNA  NM_005145 Homo sapiens gastric intrinsic factor (vitamin B synthesis) (GIF), mRNA  NM_005104 NM_005110 Homo sapiens glutamine-fructose-6-phosphate transaminase 2 (GFPT2), mRNA  NM_005104 NM_005105 Homo sapiens fusion, derived from t(12;16) malignant liposarcoma (FUS), mRNA  NM_004969 Homo sapiens nuclear receptor subfamily 5, group A, member 1 (NR5A1), mRNA		3 (KCND3), mRNA
NM_004978 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 4 (KCNC4), mRNA  NM_004977 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 3 (KCNC3), mRNA  NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNC1), mRNA  NM_004975 Homo sapiens potassium voltage-gated channel, Shab-related subfamily, member 1 (KCNB1), mRNA  NM_004969 Homo sapiens insulin-degrading enzyme (IDE), mRNA  NM_005143 Homo sapiens haptoglobin (HP), mRNA  NM_005144 Homo sapiens high-mobility group (nonhistone chromosomal) protein 14 (HMG14), mRNA  NM_005130 Homo sapiens heparin-binding growth factor binding protein (HBP17), mRNA  NM_004963 Homo sapiens guanylate cyclase 2C (heat stable enterotoxin receptor) (GUCY2C), mRNA  NM_005100 Homo sapiens A kinase (PRKA) anchor protein (gravin) 12 (AKAP12), mRNA  NM_005113 Homo sapiens golgi autoantigen, golgin subfamily a, 5 (GOLGA5), mRNA  NM_005145 Homo sapiens guanine nucleotide binding protein (G protein), gamma 7 (GNG7), mRNA  NM_005142 Homo sapiens gastric intrinsic factor (vitamin B synthesis) (GIF), mRNA  NM_00510 Homo sapiens glutamine-fructose-6-phosphate transaminase 2 (GFPT2), mRNA  NM_00510 Homo sapiens fusion, derived from t(12;16) malignant liposarcoma (FUS), mRNA  NM_004969 Homo sapiens nuclear receptor subfamily 5, group A, member 1 (NR5A1), mRNA	NM_004979	
member 4 (KCNC4), mRNA  NM_004977 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 3 (KCNC3), mRNA  NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNC1), mRNA  NM_004975 Homo sapiens potassium voltage-gated channel, Shab-related subfamily, member 1 (KCNB1), mRNA  NM_004969 Homo sapiens insulin-degrading enzyme (IDE), mRNA  NM_005143 Homo sapiens haptoglobin (HP), mRNA  NM_004965 Homo sapiens high-mobility group (nonhistone chromosomal) protein 14 (HMG14), mRNA  NM_005130 Homo sapiens heparin-binding growth factor binding protein (HBP17), mRNA  NM_004963 Homo sapiens guanylate cyclase 2C (heat stable enterotoxin receptor) (GUCY2C), mRNA  NM_005100 Homo sapiens A kinase (PRKA) anchor protein (gravin) 12 (AKAP12), mRNA  NM_005113 Homo sapiens golgi autoantigen, golgin subfamily a, 5 (GOLGA5), mRNA  NM_005145 Homo sapiens guanine nucleotide binding protein (G protein), gamma 7 (GNG7), mRNA  NM_005142 Homo sapiens gastric intrinsic factor (vitamin B synthesis) (GIF), mRNA  NM_005140 Homo sapiens glutamine-fructose-6-phosphate transaminase 2 (GFPT2), mRNA  NM_005141 Homo sapiens fusion, derived from t(12;16) malignant liposarcoma (FUS), mRNA  NM_004969 Homo sapiens nuclear receptor subfamily 5, group A, member 1 (NR5A1), mRNA		(KCND1), mRNA
NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 3 (KCNC3), mRNA  NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNC1), mRNA  NM_004975 Homo sapiens potassium voltage-gated channel, Shab-related subfamily, member 1 (KCNB1), mRNA  NM_004969 Homo sapiens insulin-degrading enzyme (IDE), mRNA  NM_005143 Homo sapiens haptoglobin (HP), mRNA  NM_004965 Homo sapiens high-mobility group (nonhistone chromosomal) protein 14 (HMG14), mRNA  NM_005130 Homo sapiens heparin-binding growth factor binding protein (HBP17), mRNA  NM_004963 Homo sapiens guanylate cyclase 2C (heat stable enterotoxin receptor) (GUCY2C), mRNA  NM_005100 Homo sapiens A kinase (PRKA) anchor protein (gravin) 12 (AKAP12), mRNA  NM_005113 Homo sapiens golgi autoantigen, golgin subfamily a, 5 (GOLGA5), mRNA  NM_005145 Homo sapiens guanine nucleotide binding protein (G protein), gamma 7 (GNG7), mRNA  NM_005142 Homo sapiens gastric intrinsic factor (vitamin B synthesis) (GIF), mRNA  NM_00510 Homo sapiens glutamine-fructose-6-phosphate transaminase 2 (GFPT2), mRNA  NM_004960 Homo sapiens fusion, derived from t(12;16) malignant liposarcoma (FUS), mRNA  NM_004959 Homo sapiens nuclear receptor subfamily 5, group A, member 1 (NR5A1), mRNA	NM_004978	Homo sapiens potassium voltage-gated channel, Shaw-related subfamily,
member 3 (KCNC3), mRNA  NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNC1), mRNA  NM_004975 Homo sapiens potassium voltage-gated channel, Shab-related subfamily, member 1 (KCNB1), mRNA  NM_004969 Homo sapiens insulin-degrading enzyme (IDE), mRNA  NM_005143 Homo sapiens haptoglobin (HP), mRNA  NM_004965 Homo sapiens high-mobility group (nonhistone chromosomal) protein 14 (HMG14), mRNA  NM_005130 Homo sapiens heparin-binding growth factor binding protein (HBP17), mRNA  NM_004963 Homo sapiens guanylate cyclase 2C (heat stable enterotoxin receptor) (GUCY2C), mRNA  NM_005100 Homo sapiens A kinase (PRKA) anchor protein (gravin) 12 (AKAP12), mRNA  NM_005113 Homo sapiens golgi autoantigen, golgin subfamily a, 5 (GOLGA5), mRNA  NM_005145 Homo sapiens guanine nucleotide binding protein (G protein), gamma 7 (GNG7), mRNA  NM_005140 Homo sapiens gastric intrinsic factor (vitamin B synthesis) (GIF), mRNA  NM_00510 Homo sapiens glutamine-fructose-6-phosphate transaminase 2 (GFPT2), mRNA  NM_004960 Homo sapiens fusion, derived from t(12;16) malignant liposarcoma (FUS), mRNA  NM_004959 Homo sapiens nuclear receptor subfamily 5, group A, member 1 (NR5A1), mRNA		member 4 (KCNC4), mRNA
NM_004976 Homo sapiens potassium voltage-gated channel, Shaw-related subfamily, member 1 (KCNC1), mRNA  NM_004975 Homo sapiens potassium voltage-gated channel, Shab-related subfamily, member 1 (KCNB1), mRNA  NM_004969 Homo sapiens insulin-degrading enzyme (IDE), mRNA  NM_005143 Homo sapiens haptoglobin (HP), mRNA  NM_004965 Homo sapiens high-mobility group (nonhistone chromosomal) protein 14 (HMG14), mRNA  NM_005130 Homo sapiens heparin-binding growth factor binding protein (HBP17), mRNA  NM_004963 Homo sapiens guanylate cyclase 2C (heat stable enterotoxin receptor) (GUCY2C), mRNA  NM_005100 Homo sapiens A kinase (PRKA) anchor protein (gravin) 12 (AKAP12), mRNA  NM_005113 Homo sapiens golgi autoantigen, golgin subfamily a, 5 (GOLGA5), mRNA  NM_005145 Homo sapiens guanine nucleotide binding protein (G protein), gamma 7 (GNG7), mRNA  NM_005140 Homo sapiens gastric intrinsic factor (vitamin B synthesis) (GIF), mRNA  NM_00510 Homo sapiens glutamine-fructose-6-phosphate transaminase 2 (GFPT2), mRNA  NM_00510 Homo sapiens fusion, derived from t(12;16) malignant liposarcoma (FUS), mRNA  NM_004960 Homo sapiens nuclear receptor subfamily 5, group A, member 1 (NR5A1), mRNA	NM_004977	Homo sapiens potassium voltage-gated channel, Shaw-related subfamily,
member 1 (KCNC1), mRNA  NM_004975 Homo sapiens potassium voltage-gated channel, Shab-related subfamily, member 1 (KCNB1), mRNA  NM_004969 Homo sapiens insulin-degrading enzyme (IDE), mRNA  NM_005143 Homo sapiens haptoglobin (HP), mRNA  NM_004965 Homo sapiens high-mobility group (nonhistone chromosomal) protein 14 (HMG14), mRNA  NM_005130 Homo sapiens heparin-binding growth factor binding protein (HBP17), mRNA  NM_004963 Homo sapiens guanylate cyclase 2C (heat stable enterotoxin receptor) (GUCY2C), mRNA  NM_005100 Homo sapiens A kinase (PRKA) anchor protein (gravin) 12 (AKAP12), mRNA  NM_005113 Homo sapiens golgi autoantigen, golgin subfamily a, 5 (GOLGA5), mRNA  NM_005145 Homo sapiens guanine nucleotide binding protein (G protein), gamma 7 (GNG7), mRNA  NM_005142 Homo sapiens gastric intrinsic factor (vitamin B synthesis) (GIF), mRNA  NM_005110 Homo sapiens glutamine-fructose-6-phosphate transaminase 2 (GFPT2), mRNA  NM_004960 Homo sapiens fusion, derived from t(12;16) malignant liposarcoma (FUS), mRNA  NM_004959 Homo sapiens nuclear receptor subfamily 5, group A, member 1 (NR5A1), mRNA		member 3 (KCNC3), mRNA
NM_004975 Homo sapiens potassium voltage-gated channel, Shab-related subfamily, member 1 (KCNB1), mRNA  NM_004969 Homo sapiens insulin-degrading enzyme (IDE), mRNA  NM_005143 Homo sapiens haptoglobin (HP), mRNA  NM_004965 Homo sapiens high-mobility group (nonhistone chromosomal) protein 14 (HMG14), mRNA  NM_005130 Homo sapiens heparin-binding growth factor binding protein (HBP17), mRNA  NM_004963 Homo sapiens guanylate cyclase 2C (heat stable enterotoxin receptor) (GUCY2C), mRNA  NM_005100 Homo sapiens A kinase (PRKA) anchor protein (gravin) 12 (AKAP12), mRNA  NM_005113 Homo sapiens golgi autoantigen, golgin subfamily a, 5 (GOLGA5), mRNA  NM_005145 Homo sapiens guanine nucleotide binding protein (G protein), gamma 7 (GNG7), mRNA  NM_005142 Homo sapiens gastric intrinsic factor (vitamin B synthesis) (GIF), mRNA  NM_00510 Homo sapiens glutamine-fructose-6-phosphate transaminase 2 (GFPT2), mRNA  NM_004960 Homo sapiens fusion, derived from t(12;16) malignant liposarcoma (FUS), mRNA  NM_004959 Homo sapiens nuclear receptor subfamily 5, group A, member 1 (NR5A1), mRNA	NM_004976	
NM_004969 Homo sapiens insulin-degrading enzyme (IDE), mRNA  NM_005143 Homo sapiens haptoglobin (HP), mRNA  NM_004965 Homo sapiens high-mobility group (nonhistone chromosomal) protein 14 (HMG14), mRNA  NM_005130 Homo sapiens heparin-binding growth factor binding protein (HBP17), mRNA  NM_004963 Homo sapiens guanylate cyclase 2C (heat stable enterotoxin receptor) (GUCY2C), mRNA  NM_005100 Homo sapiens A kinase (PRKA) anchor protein (gravin) 12 (AKAP12), mRNA  NM_005113 Homo sapiens golgi autoantigen, golgin subfamily a, 5 (GOLGA5), mRNA  NM_005145 Homo sapiens guanine nucleotide binding protein (G protein), gamma 7 (GNG7), mRNA  NM_005142 Homo sapiens gastric intrinsic factor (vitamin B synthesis) (GIF), mRNA  NM_005110 Homo sapiens glutamine-fructose-6-phosphate transaminase 2 (GFPT2), mRNA  NM_004960 Homo sapiens fusion, derived from t(12;16) malignant liposarcoma (FUS), mRNA  NM_004959 Homo sapiens nuclear receptor subfamily 5, group A, member 1 (NR5A1), mRNA		member 1 (KCNC1), mRNA
NM_004969 Homo sapiens insulin-degrading enzyme (IDE), mRNA  NM_005143 Homo sapiens haptoglobin (HP), mRNA  NM_004965 Homo sapiens high-mobility group (nonhistone chromosomal) protein 14 (HMG14), mRNA  NM_005130 Homo sapiens heparin-binding growth factor binding protein (HBP17), mRNA  NM_004963 Homo sapiens guanylate cyclase 2C (heat stable enterotoxin receptor) (GUCY2C), mRNA  NM_005100 Homo sapiens A kinase (PRKA) anchor protein (gravin) 12 (AKAP12), mRNA  NM_005113 Homo sapiens golgi autoantigen, golgin subfamily a, 5 (GOLGA5), mRNA  NM_005145 Homo sapiens guanine nucleotide binding protein (G protein), gamma 7 (GNG7), mRNA  NM_005142 Homo sapiens gastric intrinsic factor (vitamin B synthesis) (GIF), mRNA  NM_005110 Homo sapiens glutamine-fructose-6-phosphate transaminase 2 (GFPT2), mRNA  NM_004960 Homo sapiens fusion, derived from t(12;16) malignant liposarcoma (FUS), mRNA  NM_004959 Homo sapiens nuclear receptor subfamily 5, group A, member 1 (NR5A1), mRNA	NM_004975	Homo sapiens potassium voltage-gated channel, Shab-related subfamily, member
NM_005143 Homo sapiens haptoglobin (HP), mRNA  NM_004965 Homo sapiens high-mobility group (nonhistone chromosomal) protein 14 (HMG14), mRNA  NM_005130 Homo sapiens heparin-binding growth factor binding protein (HBP17), mRNA  NM_004963 Homo sapiens guanylate cyclase 2C (heat stable enterotoxin receptor) (GUCY2C), mRNA  NM_005100 Homo sapiens A kinase (PRKA) anchor protein (gravin) 12 (AKAP12), mRNA  NM_005113 Homo sapiens golgi autoantigen, golgin subfamily a, 5 (GOLGA5), mRNA  NM_005145 Homo sapiens guanine nucleotide binding protein (G protein), gamma 7 (GNG7), mRNA  NM_005142 Homo sapiens gastric intrinsic factor (vitamin B synthesis) (GIF), mRNA  NM_005110 Homo sapiens glutamine-fructose-6-phosphate transaminase 2 (GFPT2), mRNA  NM_004960 Homo sapiens fusion, derived from t(12;16) malignant liposarcoma (FUS), mRNA  NM_004959 Homo sapiens nuclear receptor subfamily 5, group A, member 1 (NR5A1), mRNA	_	1 (KCNB1), mRNA
NM_004965 Homo sapiens high-mobility group (nonhistone chromosomal) protein 14 (HMG14), mRNA  NM_005130 Homo sapiens heparin-binding growth factor binding protein (HBP17), mRNA  NM_004963 Homo sapiens guanylate cyclase 2C (heat stable enterotoxin receptor) (GUCY2C), mRNA  NM_005100 Homo sapiens A kinase (PRKA) anchor protein (gravin) 12 (AKAP12), mRNA  NM_005113 Homo sapiens golgi autoantigen, golgin subfamily a, 5 (GOLGA5), mRNA  NM_005145 Homo sapiens guanine nucleotide binding protein (G protein), gamma 7 (GNG7), mRNA  NM_005142 Homo sapiens gastric intrinsic factor (vitamin B synthesis) (GIF), mRNA  NM_005110 Homo sapiens glutamine-fructose-6-phosphate transaminase 2 (GFPT2), mRNA  NM_004960 Homo sapiens fusion, derived from t(12;16) malignant liposarcoma (FUS), mRNA  NM_004959 Homo sapiens nuclear receptor subfamily 5, group A, member 1 (NR5A1), mRNA	NM 004969	Homo sapiens insulin-degrading enzyme (IDE), mRNA
NM_004965 Homo sapiens high-mobility group (nonhistone chromosomal) protein 14 (HMG14), mRNA  NM_005130 Homo sapiens heparin-binding growth factor binding protein (HBP17), mRNA  NM_004963 Homo sapiens guanylate cyclase 2C (heat stable enterotoxin receptor) (GUCY2C), mRNA  NM_005100 Homo sapiens A kinase (PRKA) anchor protein (gravin) 12 (AKAP12), mRNA  NM_005113 Homo sapiens golgi autoantigen, golgin subfamily a, 5 (GOLGA5), mRNA  NM_005145 Homo sapiens guanine nucleotide binding protein (G protein), gamma 7 (GNG7), mRNA  NM_005142 Homo sapiens gastric intrinsic factor (vitamin B synthesis) (GIF), mRNA  NM_005110 Homo sapiens glutamine-fructose-6-phosphate transaminase 2 (GFPT2), mRNA  NM_004960 Homo sapiens fusion, derived from t(12;16) malignant liposarcoma (FUS), mRNA  NM_004959 Homo sapiens nuclear receptor subfamily 5, group A, member 1 (NR5A1), mRNA	NM 005143	Homo sapiens haptoglobin (HP), mRNA
(HMG14), mRNA  NM_005130 Homo sapiens heparin-binding growth factor binding protein (HBP17), mRNA  NM_004963 Homo sapiens guanylate cyclase 2C (heat stable enterotoxin receptor) (GUCY2C), mRNA  NM_005100 Homo sapiens A kinase (PRKA) anchor protein (gravin) 12 (AKAP12), mRNA  NM_005113 Homo sapiens golgi autoantigen, golgin subfamily a, 5 (GOLGA5), mRNA  NM_005145 Homo sapiens guanine nucleotide binding protein (G protein), gamma 7 (GNG7), mRNA  NM_005142 Homo sapiens gastric intrinsic factor (vitamin B synthesis) (GIF), mRNA  NM_005110 Homo sapiens glutamine-fructose-6-phosphate transaminase 2 (GFPT2), mRNA  NM_004960 Homo sapiens fusion, derived from t(12;16) malignant liposarcoma (FUS), mRNA  NM_004959 Homo sapiens nuclear receptor subfamily 5, group A, member 1 (NR5A1), mRNA		Homo sapiens high-mobility group (nonhistone chromosomal) protein 14
NM_005130 Homo sapiens heparin-binding growth factor binding protein (HBP17), mRNA  NM_004963 Homo sapiens guanylate cyclase 2C (heat stable enterotoxin receptor) (GUCY2C), mRNA  NM_005100 Homo sapiens A kinase (PRKA) anchor protein (gravin) 12 (AKAP12), mRNA  NM_005113 Homo sapiens golgi autoantigen, golgin subfamily a, 5 (GOLGA5), mRNA  NM_005145 Homo sapiens guanine nucleotide binding protein (G protein), gamma 7 (GNG7), mRNA  NM_005142 Homo sapiens gastric intrinsic factor (vitamin B synthesis) (GIF), mRNA  NM_005110 Homo sapiens glutamine-fructose-6-phosphate transaminase 2 (GFPT2), mRNA  NM_004960 Homo sapiens fusion, derived from t(12;16) malignant liposarcoma (FUS), mRNA  NM_004959 Homo sapiens nuclear receptor subfamily 5, group A, member 1 (NR5A1), mRNA		
NM_004963 Homo sapiens guanylate cyclase 2C (heat stable enterotoxin receptor) (GUCY2C), mRNA  NM_005100 Homo sapiens A kinase (PRKA) anchor protein (gravin) 12 (AKAP12), mRNA  NM_005113 Homo sapiens golgi autoantigen, golgin subfamily a, 5 (GOLGA5), mRNA  NM_005145 Homo sapiens guanine nucleotide binding protein (G protein), gamma 7 (GNG7), mRNA  NM_005142 Homo sapiens gastric intrinsic factor (vitamin B synthesis) (GIF), mRNA  NM_005110 Homo sapiens glutamine-fructose-6-phosphate transaminase 2 (GFPT2), mRNA  NM_004960 Homo sapiens fusion, derived from t(12;16) malignant liposarcoma (FUS), mRNA  NM_004959 Homo sapiens nuclear receptor subfamily 5, group A, member 1 (NR5A1), mRNA	NM 005130	Homo sapiens heparin-binding growth factor binding protein (HBP17), mRNA
(GUCY2C), mRNA  NM 005100 Homo sapiens A kinase (PRKA) anchor protein (gravin) 12 (AKAP12), mRNA  NM 005113 Homo sapiens golgi autoantigen, golgin subfamily a, 5 (GOLGA5), mRNA  NM 005145 Homo sapiens guanine nucleotide binding protein (G protein), gamma 7  (GNG7), mRNA  NM 005142 Homo sapiens gastric intrinsic factor (vitamin B synthesis) (GIF), mRNA  NM 005110 Homo sapiens glutamine-fructose-6-phosphate transaminase 2 (GFPT2), mRNA  NM 004960 Homo sapiens fusion, derived from t(12;16) malignant liposarcoma (FUS),  mRNA  NM 004959 Homo sapiens nuclear receptor subfamily 5, group A, member 1 (NR5A1),  mRNA		Homo sapiens guanylate cyclase 2C (heat stable enterotoxin receptor)
NM_005100 Homo sapiens A kinase (PRKA) anchor protein (gravin) 12 (AKAP12), mRNA  NM_005113 Homo sapiens golgi autoantigen, golgin subfamily a, 5 (GOLGA5), mRNA  NM_005145 Homo sapiens guanine nucleotide binding protein (G protein), gamma 7  (GNG7), mRNA  NM_005142 Homo sapiens gastric intrinsic factor (vitamin B synthesis) (GIF), mRNA  NM_005110 Homo sapiens glutamine-fructose-6-phosphate transaminase 2 (GFPT2), mRNA  NM_004960 Homo sapiens fusion, derived from t(12;16) malignant liposarcoma (FUS),  mRNA  NM_004959 Homo sapiens nuclear receptor subfamily 5, group A, member 1 (NR5A1),  mRNA		
NM_005113 Homo sapiens golgi autoantigen, golgin subfamily a, 5 (GOLGA5), mRNA  NM_005145 Homo sapiens guanine nucleotide binding protein (G protein), gamma 7 (GNG7), mRNA  NM_005142 Homo sapiens gastric intrinsic factor (vitamin B synthesis) (GIF), mRNA  NM_005110 Homo sapiens glutamine-fructose-6-phosphate transaminase 2 (GFPT2), mRNA  NM_004960 Homo sapiens fusion, derived from t(12;16) malignant liposarcoma (FUS), mRNA  NM_004959 Homo sapiens nuclear receptor subfamily 5, group A, member 1 (NR5A1), mRNA	NM 005100	Homo saniens A kinase (PRKA) anchor protein (gravin) 12 (AKAP12), mRNA
NM_005145 Homo sapiens guanine nucleotide binding protein (G protein), gamma 7 (GNG7), mRNA  NM_005142 Homo sapiens gastric intrinsic factor (vitamin B synthesis) (GIF), mRNA  NM_005110 Homo sapiens glutamine-fructose-6-phosphate transaminase 2 (GFPT2), mRNA  NM_004960 Homo sapiens fusion, derived from t(12;16) malignant liposarcoma (FUS), mRNA  NM_004959 Homo sapiens nuclear receptor subfamily 5, group A, member 1 (NR5A1), mRNA		Homo saniens golgi autoantigen, golgin subfamily a. 5 (GOLGA5), mRNA
(GNG7), mRNA  NM_005142 Homo sapiens gastric intrinsic factor (vitamin B synthesis) (GIF), mRNA  NM_005110 Homo sapiens glutamine-fructose-6-phosphate transaminase 2 (GFPT2), mRNA  NM_004960 Homo sapiens fusion, derived from t(12;16) malignant liposarcoma (FUS),  mRNA  NM_004959 Homo sapiens nuclear receptor subfamily 5, group A, member 1 (NR5A1),  mRNA		Homo sapiens guarine nucleotide hinding protein (G protein) gamma 7
NM_005142 Homo sapiens gastric intrinsic factor (vitamin B synthesis) (GIF), mRNA  NM_005110 Homo sapiens glutamine-fructose-6-phosphate transaminase 2 (GFPT2), mRNA  NM_004960 Homo sapiens fusion, derived from t(12;16) malignant liposarcoma (FUS),  mRNA  NM_004959 Homo sapiens nuclear receptor subfamily 5, group A, member 1 (NR5A1),  mRNA	14147_002142	
NM_005110 Homo sapiens glutamine-fructose-6-phosphate transaminase 2 (GFPT2), mRNA  NM_004960 Homo sapiens fusion, derived from t(12;16) malignant liposarcoma (FUS), mRNA  NM_004959 Homo sapiens nuclear receptor subfamily 5, group A, member 1 (NR5A1), mRNA	NTM 005142	
NM_004960 Homo sapiens fusion, derived from t(12;16) malignant liposarcoma (FUS), mRNA  NM_004959 Homo sapiens nuclear receptor subfamily 5, group A, member 1 (NR5A1), mRNA		The residue electronic fraction (vitalian B synthesis) (Gr.), madva
mRNA NM_004959 Homo sapiens nuclear receptor subfamily 5, group A, member 1 (NR5A1), mRNA		Homo sapiens giutamine-iruciose-o-phosphate transaminase 2 (GPT 12), likiva
NM_004959 Homo sapiens nuclear receptor subfamily 5, group A, member 1 (NR5A1), mRNA	NM_004960	•
mRNA		mkna
	NM_004959	•
NM_004957 Homo sapiens folylpolyglutamate synthase (FPGS), mRNA		
	NM_004957	Homo sapiens folylpolyglutamate synthase (FPGS), mRNA

. WO 03/074654

7. 004056	Homo sapiens ets variant gene 1 (ETV1), mRNA
VM_004956	Homo sapiens ets variant gene 1 (ETV1), micric Homo sapiens solute carrier family 29 (nucleoside transporters), member 1
	- 1 C 1:1 1 (LNU)C)CT
NM 005107	Homo sapiens endonuclease G-like 1 (ENDOGET), indexing Homo sapiens eukaryotic translation initiation factor 4 gamma, 1 (EIF4G1),
NM_004953	mRNA
27.6.004052	in a colorin A2 (EENA3) mRNA
NM_004952	1
NM_004944	
NM_004938	
NM_005127	1 discontinuity member / (activation-induced) (CEE-CEE)
NT 6 004025	1' 1donf langee 3 11 .1 N. J. 111111111
NM_004935	
NM_004931	
NM_005125	Homo sapiens core-binding factor, runt domain, alpha subunit 2; translocated to,
NM_005093	2 (CBFA2T2), mRNA
NIM 004020	2 (CBFA2T2), mRNA  Homo sapiens capping protein (actin filament) muscle Z-line, beta (CAPZB),
NM_004930	mP.NA
ND 6 005120	· A 2 (ANTV A 3) mRNA
NM_005139	Home caniens acetyl-Coenzyme A carboxylase aipha (11011011);
NM_000664	Transport dine ammonia-IVASE I II/AL/, IIII/1/12
NM_002108	Homo sapiens bone morphogenetic protein 6 (BMP6), mRNA
NM_001718	Homo sapiens annexin A5 (ANXA5), mRNA  Homo sapiens annexin A5 (ANXA5), mRNA
NM_001154	
NM_001153	
NM_004817	
NM_004736	Homo sapiens xeroderma pigmentosum, complementation group C (XPC),
NM_004628	DATA
	mRNA Homo sapiens tryptophan rich basic protein (WRB), mRNA (Magt disease, bestrophin) (VMD2)
NM_004627	Homo sapiens tryptophan rich basic protein (WRD), interval Homo sapiens vitelliform macular dystrophy (Best disease, bestrophin) (VMD2)
NM_004183	
	mRNA Homo sapiens Vertebrate LIN7 homolog 1, Tax interaction protein 33 (VELI1),
NM_004664	DATA
	mRNA  Homo sapiens variable charge, Y chromosome (VCY), mRNA
NM_004679	- Lar whigh touch expressed transcript (UAT), interest
NM_004182	
NM_004651	Homo sapiens ubiquitin specific protease 11 (OSI 17), mediate Homo sapiens ubiquitin carboxyl-terminal esterase L1 (ubiquitin thiolesterase)
NM_004181	
	TI which conjugating enzyme E2L 6 (UBE2LO), HIGHA
NM_004223	
NM_004623	
NM_004622	the world recentor interacting biological 15 (1101 15); many
NM_004236	Try torrel registance associated gene 5 (110105), ind to
NM_004909	
NM_004295	
NM_004179	D374
	mRNA Homo sapiens tumor necrosis factor receptor superfamily, member 18
NM_004195	
	(TNFRSF18), mRNA  Homo sapiens thymosin, beta 4, Y chromosome (TMSB4Y), mRNA  (TM4SF3), mRNA
NM_004202	
NM_00461	
NM_00461	
NM 00486	5 Homo sapiens TBP-like 1 (TBPL1), mRNA

Homo sapiens transglutaminase 2 (C polypeptide, protein-glutamine-gamma-glutamyltransferase) (TGM2), mRNA
Homo sapiens transforming growth factor, beta receptor I (activin A receptor
type II-like kinase, 53kD) (TGFBR1), mRNA
Homo sapiens programmed cell death 5 (PDCD5), mRNA
Homo sapiens T-cell leukemia/lymphoma 1B (TCL1B), mRNA
Homo sapiens transcription factor 15 (basic helix-loop-helix) (TCF15), mRNA
Homo sapiens transcription elongation factor A (SII)-like 1 (TCEAL1), mRNA
Homo sapiens thousand and one amino acid protein kinase (TAO1), mRNA
Homo sapiens TATA box binding protein (TBP)-associated factor, RNA
polymerase II, A, 250kD (TAF2A), mRNA
Homo sapiens synaptogyrin 2 (SYNGR2), mRNA
Homo sapiens synaptogyrin 1 (SYNGR1), mRNA
Homo sapiens sulfotransferase family, cytosolic, 2B, member 1 (SULT2B1), mRNA
Homo sapiens syntaxin 8 (STX8), mRNA
Homo sapiens syntaxin 1A (brain) (STX1A), mRNA
Homo sapiens serine/threonine kinase 12 (STK12), mRNA
Homo sapiens sterol regulatory element binding transcription factor 2 (SREBF2),
mRNA
Homo sapiens sterol regulatory element binding transcription factor 1 (SREBF1), mRNA
Homo sapiens secreted phosphoprotein 1 (osteopontin, bone sialoprotein I, early
T-lymphocyte activation 1) (SPP1), mRNA
Homo sapiens SRY (sex determining region Y)-box 14 (SOX14), mRNA
Homo sapiens small nuclear ribonucleoprotein polypeptide A (SNRPA), mRNA
Homo sapiens synaptosomal-associated protein, 29kD (SNAP29), mRNA
Homo sapiens spermine synthase (SMS), mRNA
Homo sapiens solute carrier family 9 (sodium/hydrogen exchanger), isoform 5 (SLC9A5), mRNA
Homo sapiens solute carrier family 7 (cationic amino acid transporter, y+system), member 4 (SLC7A4), mRNA
Homo sapiens solute carrier family 6 (neurotransmitter transporter, glycine), member 5 (SLC6A5), mRNA
Homo sapiens solute carrier family 4, sodium bicarbonate cotransporter, member 8 (SLC4A8), mRNA
Homo sapiens solute carrier family 24 (sodium/potassium/calcium exchanger), member 1 (SLC24A1), mRNA
Homo sapiens solute carrier family 1 (glial high affinity glutamate transporter), member 3 (SLC1A3), nuclear gene encoding mitochondrial protein, mRNA
Homo sapiens solute carrier family 1 (glial high affinity glutamate transporter),
member 2 (SLC1A2), nuclear gene encoding mitochondrial protein, mRNA
Homo sapiens solute carrier family 16 (monocarboxylic acid transporters),
member 7 (SLC16A7), mRNA
Homo sapiens solute carrier family 16 (monocarboxylic acid transporters),
member 5 (SLC16A5), mRNA  Homo sapiens solute carrier family 16 (monocarboxylic acid transporters),
member 3 (SLC16A3), mRNA
Homo sapiens mannose-P-dolichol utilization defect 1 (MPDU1), mRNA
Homo sapiens splicing factor, arginine/serine-rich 11 (SFRS11), mRNA
Homo sapiens sema domain, immunoglobulin domain (Ig), short basic domain,

	1 (GDD1) DVA
NM_004753	Homo sapiens short-chain dehydrogenase/reductase 1 (SDR1), mRNA
NM 004168	Homo sapiens succinate dehydrogenase complex, subunit A, flavoprotein (Fp)
_	(SDHA), nuclear gene encoding mitochondrial protein, mRNA
NM 004713	Homo saniens serologically defined colon cancer antigen 1 (SDCCAG1), HIGNA
NM_004591	Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 20
_	(SCYA20) mRNA
NM_004590	Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 16
	(SCYA16), mRNA
NM_004588	Homo sapiens sodium channel, voltage-gated, type II, beta polypeptide
1111_00 1000	(SCN2B), mRNA
NM_004165	Homo sapiens Ras-related associated with diabetes (RRAD), mRNA
NM_004755	Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 5 (RPS6KA5),
14141_00-1755	mRNA
NM_004586	Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 3 (RPS6KA3),
14141_004200	mRNA
NM_004790	Homo sapiens solute carrier family 22 (organic anion transporter), member 6
MM_004790	(SLC22A6), mRNA
NM 004259	Homo sapiens RecQ protein-like 5 (RECQL5), mRNA
	Homo sapiens RecQ protein-like 4 (RECQL4), mRNA
NM_004260	Homo sapiens RAB5C, member RAS oncogene family (RAB5C), mRNA
NM_004583	Homo sapiens Rab geranylgeranyltransferase, beta subunit (RABGGTB), mRNA
NM_004582	Homo sapiens Rab geranylgeranyltransferase, alpha subunit (RABGGTA),
NM_004581	
	mRNA Homo sapiens RAB9, member RAS oncogene family (RAB9), mRNA
NM_004251	Homo sapiens RAB9, member RAS officogene family (RAB5A) mRNA
NM_004162	Homo sapiens RAB5A, member RAS oncogene family (RAB5A), mRNA
NM_004578	Homo sapiens RAB4, member RAS oncogene family (RAB4), mRNA
NM_004914	Homo sapiens RAB36, member RAS oncogene family (RAB36), mRNA
NM_004580	Homo sapiens RAB27A, member RAS oncogene family (RAB27A), mRNA
NM_004663	Homo sapiens RAB11A, member RAS oncogene family (RAB11A), mRNA
NM_004160	Homo sapiens peptide YY (PYY), mRNA
NM_004103	Homo sapiens protein tyrosine kinase 2 beta (PTK2B), mRNA
NM_004158	Homo sapiens persephin (PSPN), mRNA
NM 004577	Homo sapiens phosphoserine phosphatase (PSPH), mRNA
NM_004159	Homo sapiens proteasome (prosome, macropain) subunit, beta type, 8 (large
	multifunctional protease 7) (PSMB8), mRNA
NM 004917	Homo sapiens kallikrein 4 (prostase, enamel matrix, prostate) (KLK4), mRNA
NM_004157	Homo sapiens protein kinase, cAMP-dependent, regulatory, type II, alpha
_	(PRKAR2A), mRNA
NM_004758	Homo sapiens peripheral benzodiazepine receptor-associated protein 1 (PRAX-
1,1,1	1). mRNA
NM 004576	Homo sapiens protein phosphatase 2 (formerly 2A), regulatory subunit B (PR
14142_00 10 10	52), beta isoform (PPP2R2B), mRNA
NM_004156	Homo sapiens protein phosphatase 2 (formerly 2A), catalytic subunit, beta
14141_00 1150	isoform (PPP2CB), mRNA
NM 000942	Homo sapiens peptidylprolyl isomerase B (cyclophilin B) (PPIB), mRNA
NM_004575	Homo sapiens POU domain, class 4, transcription factor 2 (POU4F2), mRNA
NM 004573	Homo sapiens phospholipase C, beta 2 (PLCB2), mRNA
NM 004572	Homo sapiens plakophilin 2 (PKP2), mRNA
	Homo sapiens PBX/knotted 1 hoemobox 1 (PKNOX1), mRNA
NM_004571	Homo sapiens membrane-associated tyrosine- and threonine-specific cdc2-
NM_004203	inhibitory kinase (PKMYT1), mRNA
NR 5 004010	Homo sapiens phosphatidylinositol transfer protein, membrane-associated
NM_004910	nomo sapiens phosphaticy iniciation in ansier protein, memorane-associated

<del></del> 1	(DITDNIM)DNIA
ND 4 00 4279	(PITPNM), mRNA  Homo sapiens phosphatidylinositol glycan, class L (PIGL), mRNA
NM_004278	Homo sapiens phosphatidylinositol glycan, class E (FIGE), mRNA  Homo sapiens phosphatidylinositol glycan, class H (PIGH), mRNA
NM_004569	Homo sapiens phosphatidylmositol glycan, class B (PIGB), mRNA
NM_004855	Homo sapiens LPS-induced TNF-alpha factor (PIG7), mRNA
NM_004862	Homo sapiens prostaglandin E synthase (PTGES), mRNA
NM_004878	Homo sapiens 6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 4
NM_004567	(PFKFB4), mRNA
NM_004566	Homo sapiens 6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 3 (PFKFB3), mRNA
NM_004836	Homo sapiens eukaryotic translation initiation factor 2-alpha kinase 3 (EIF2AK3), mRNA
NM 004716	Homo sapiens proprotein convertase subtilisin/kexin type 7 (PCSK7), mRNA
NM_000437	Homo sapiens platelet-activating factor acetylhydrolase 2 (40kD) (PAFAH2), mRNA
NM_004199	Homo sapiens procollagen-proline, 2-oxoglutarate 4-dioxygenase (proline 4-hydroxylase), alpha polypeptide II (P4HA2), mRNA
NM_004154	Homo sapiens pyrimidinergic receptor P2Y, G-protein coupled, 6 (P2RY6), mRNA
NM_004280	Homo sapiens eukaryotic translation elongation factor 1 epsilon 1 (EEF1E1), mRNA
NM 004741	Homo sapiens nucleolar phosphoprotein p130 (P130), mRNA
NM 004802	Homo sapiens otoferlin (OTOF), mRNA
NM 004852	Homo sapiens one cut domain, family member 2 (ONECUT2), mRNA
NM_004254	Homo sapiens solute carrier family 22 (organic anion transporter), member 8 (SLC22A8), mRNA
NM 004298	Homo sapiens nucleoporin 155kD (NUP155), mRNA
NM 004560	Homo sapiens receptor tyrosine kinase-like orphan receptor 2 (ROR2), mRNA
NM 004822	Homo sapiens netrin 1 (NTN1), mRNA
NM 004796	Homo sapiens neurexin 3 (NRXN3), mRNA
NM 004558	Homo sapiens neurturin (NRTN), mRNA
NM 004688	Homo sapiens N-myc (and STAT) interactor (NMI), mRNA
NM 004148	Homo sapiens ninjurin 1 (NINJ1), mRNA
NM_004552	Homo sapiens NADH dehydrogenase (ubiquinone) Fe-S protein 5 (15kD) (NADH-coenzyme Q reductase) (NDUFS5), mRNA
NM_004551	Homo sapiens NADH dehydrogenase (ubiquinone) Fe-S protein 3 (30kD) (NADH-coenzyme Q reductase) (NDUFS3), mRNA
NM_004550	Homo sapiens NADH dehydrogenase (ubiquinone) Fe-S protein 2 (49kD) (NADH-coenzyme Q reductase) (NDUFS2), mRNA
NM 004540	Homo sapiens neural cell adhesion molecule 2 (NCAM2), mRNA
NM_004644	Homo sapiens adaptor-related protein complex 3, beta 2 subunit (AP3B2), mRNA
NM 004538	Homo sapiens nucleosome assembly protein 1-like 3 (NAP1L3), mRNA
NM 004145	Homo sapiens myosin IXB (MYO9B), mRNA
NM 004294	Homo sapiens mitochondrial translational release factor 1 (MTRF1), mRNA
NM 004923	Homo sapiens metallothionein-like 5, testis-specific (tesmin) (MTL5), mRNA
NM_004143	Homo sapiens Cbp/p300-interacting transactivator, with Glu/Asp-rich carboxy-
	terminal domain, 1 (CITED1), mRNA
NM_004279	Homo sapiens peptidase (mitochondrial processing) beta (PMPCB), mRNA
NM_004531	Homo sapiens molybdenum cofactor synthesis 2 (MOCS2), mRNA
NM_004244	Homo sapiens CD163 antigen (CD163), mRNA
NM_004528	Homo sapiens microsomal glutathione S-transferase 3 (MGST3), mRNA

ND 4 004225	Homo sapiens MFH-amplified sequences with leucine-rich tandem repeats 1
NM_004225	(MASI 1) mRNA
NM 002372	Homo saniens mannosidase, alpha, class 2A, member 1 (MAN2A1), mRNA
NM_004721	Homo sapiens mitogen-activated protein kinase kinase kinase 13 (MAP3K13),
111,1	mR NA
NM_002332	Homo sapiens low density lipoprotein-related protein 1 (alpha-2-macroglobulin
1111	receptor) (LRP1), mRNA
NM 004793	Homo sapiens protease, serine, 15 (PRSS15), mRNA
NM 004789	Homo saniens LIM homeobox protein 2 (LHX2), mRNA
NM 004863	Homo sapiens serine palmitoyltransferase, long chain base subunit 2 (SPTLC2),
1111_00.000	mRNA
NM 004737	Homo sapiens like-glycosyltransferase (LARGE), mRNA
NM 004795	Homo sapiens klotho (KL), mRNA
NM 004521	Homo sapiens kinesin family member 5B (KIF5B), mRNA
NM 004520	Homo sapiens kinesin heavy chain member 2 (KIF2), mRNA
NM_004920	Homo saniens apontosis-associated tyrosine kinase (AATK), mRNA
NM 004700	Homo sapiens potassium voltage-gated channel, KQT-like subfamily, member 4
14141_004700	(KCNO4), mRNA
NM_004519	Homo sapiens potassium voltage-gated channel, KQT-like subfamily, member 3
14141_004515	(KCNQ3), mRNA
NM_004518	Homo sapiens potassium voltage-gated channel, KQT-like subfamily, member 2
14141_00-1510	(KCNQ2), mRNA
NM 004137	Homo sapiens potassium large conductance calcium-activated channel,
14141_004157	subfamily M beta member 1 (KCNMB1), mRNA
NM 004732	Homo sapiens potassium voltage-gated channel, shaker-related subfamily, beta
1414_00 1752	member 3 (KCNAB3), mRNA
NM 004693	Homo saniens cytokeratin type II (K6HF), mRNA
NM 004791	Homo sapiens integrin, beta-like 1 (with EGF-like repeat domains) (ITGBL1),
11112_00 1751	mRNA
NM 004517	Homo sapiens integrin-linked kinase (ILK), mRNA
NM 004514	Homo sapiens interleukin enhancer binding factor 1 (ILF1), mRNA
NM 004633	Homo sapiens interleukin 1 receptor, type II (IL1R2), mRNA
NM 004513	Homo sapiens interleukin 16 (lymphocyte chemoattractant factor) (IL16), mRNA
NM 004512	Homo sapiens interleukin 11 receptor, alpha (IL11RA), mRNA
NM 004258	Homo sapiens immunoglobulin superfamily, member 2 (IGSF2), mRNA
NM 004135	Homo saniens isocitrate dehydrogenase 3 (NAD+) gamma (IDH3G), mRNA
NM 004134	Homo sapiens heat shock 70kD protein 9B (mortalin-2) (HSPA9B), mRNA
NM 004697	Homo sapiens PRP4/STK/WD splicing factor (HPRP4P), mRNA
NM 004698	Homo sapiens U4/U6-associated RNA splicing factor (HPRP3P), mRNA
NM 004503	Homo sapiens homeo box C6 (HOXC6), mRNA
NM 004502	Homo sapiens homeo box B7 (HOXB7), mRNA
NM 004497	Homo sapiens hepatocyte nuclear factor 3, gamma (HNF3G), mRNA
NM 004496	Homo sapiens hepatocyte nuclear factor 3, alpha (HNF3A), mRNA
NM 004712	Homo sapiens hepatocyte growth factor-regulated tyrosine kinase substrate
14141_004/12	(HGS), mRNA
NM 004834	Homo sapiens mitogen-activated protein kinase kinase kinase kinase 4
14147_004024	(MAP4K4), mRNA
NM 004494	Homo sapiens hepatoma-derived growth factor (high-mobility group protein 1-
14141_004434	like) (HDGF), mRNA
NM 004876	Homo sapiens zinc finger protein 254 (ZNF254), mRNA
	Homo sapiens hydroxyacyl-Coenzyme A dehydrogenase, type II (HADH2),
NM_004493	mRNA
	HINNA

NIM 004004	Homo sapiens cAMP response element-binding protein CRE-BPa
NM_004904	(H GS165L15.1), mRNA
NM 004893	Homo sapiens H2A histone family, member Y (H2AFY), mRNA
NM 004130	Homo sapiens glycogenin (GYG), mRNA
NM 004286	Homo sapiens GTP binding protein 1 (GTPBP1), mRNA
NM 004128	Homo sapiens general transcription factor IIF, polypeptide 2 (30kD subunit)
1411_004120	(GTF2F2), mRNA
NM 004491	Homo sapiens glucocorticoid receptor DNA binding factor 1 (GRLF1), mRNA
NM 000826	Homo sapiens glutamate receptor, ionotropic, AMPA 2 (GRIA2), mRNA
NM 004490	Homo sapiens growth factor receptor-bound protein 14 (GRB14), mRNA
NM 004810	Homo sapiens GRB2-related adaptor protein 2 (GRAP2), mRNA
NM 004224	Homo sapiens G protein-coupled receptor 50 (GPR50), mRNA
NM 004871	Homo sapiens golgi SNAP receptor complex member 1 (GOSR1), mRNA
NM 004487	Homo sapiens golgi autoantigen, golgin subfamily b, macrogolgin (with
	transmembrane signal), 1 (GOLGB1), mRNA
NM 004126	Homo sapiens guanine nucleotide binding protein 11 (GNG11), mRNA
NM 004297	Homo sapiens guanine nucleotide binding protein (G protein), alpha 14
_	(GNA14), mRNA
NM 004246	Homo sapiens glucagon-like peptide 2 receptor (GLP2R), mRNA
NM_004123	Homo sapiens gastric inhibitory polypeptide (GIP), mRNA
NM_004121	Homo sapiens gamma-glutamyltransferase-like activity 1 (GGTLA1), mRNA
NM_004837	Homo sapiens geranylgeranyl diphosphate synthase 1 (GGPS1), mRNA
NM_004188	Homo sapiens growth factor independent 1B (potential regulator of CDKN1A,
	translocated in CML) (GFI1B), mRNA
NM_004293	Homo sapiens guanine deaminase (GDA), mRNA
NM_004751	Homo sapiens glucosaminyl (N-acetyl) transferase 3, mucin type (GCNT3), mRNA
NM 004193	Homo sapiens golgi-specific brefeldin A resistance factor 1 (GBF1), mRNA
NM 002030	Homo sapiens formyl peptide receptor-like 2 (FPRL2), mRNA
NM 004476	Homo sapiens folate hydrolase (prostate-specific membrane antigen) 1 (FOLH1),
	mRNA
NM 004119	Homo sapiens fms-related tyrosine kinase 3 (FLT3), mRNA
NM 004475	Homo sapiens flotillin 2 (FLOT2), mRNA
NM 004472	Homo sapiens forkhead box D1 (FOXD1), mRNA
NM_004471	Homo sapiens forkhead box G1A (FOXG1A), mRNA
NM_004474	Homo sapiens forkhead box D2 (FOXD2), mRNA
NM_004469	Homo sapiens c-fos induced growth factor (vascular endothelial growth factor D) (FIGF), mRNA
NM 004468	Homo sapiens four and a half LIM domains 3 (FHL3), mRNA
NM 004462	Homo sapiens four and a nari Envi domains 3 (TTE5), indexer.  Homo sapiens farnesyl-diphosphate farnesyltransferase 1 (FDFT1), mRNA
NM 004107	Homo sapiens Fc fragment of IgG, receptor, transporter, alpha (FCGRT), mRNA
NM 004107	Homo sapiens fatty acid synthase (FASN), mRNA
NM 004104 NM 004461	Homo sapiens phenylalanine-tRNA synthetase-like (FARSL), mRNA
NM 004101	Homo sapiens coagulation factor II (thrombin) receptor-like 2 (F2RL2), mRNA
NM 004101	Homo sapiens Kruppel-like factor 4 (gut) (KLF4), mRNA
NM 004455	Homo sapiens exostoses (multiple)-like 1 (EXTL1), mRNA
NM 004454	Homo sapiens ets variant gene 5 (ets-related molecule) (ETV5), mRNA
NM 004453	Homo sapiens electron-transferring-flavoprotein dehydrogenase (ETFDH),
14141_00++22	nuclear gene encoding mitochondrial protein, mRNA
NM_004452	Homo sapiens estrogen-related receptor beta (ESRRB), mRNA
	Homo sapiens protein disulfide isomerase related protein (calcium-binding
14147_00-4211	
NM_004911	Homo sapiens protein disulfide isomerase related protein (calcium-binding protein, intestinal-related) (ERP70), mRNA

	Homo sapiens epidermal growth factor receptor pathway substrate 8 (EPS8),
	70.3 T.A
7.5.00.1146	mRNA Homo sapiens glutamyl-prolyl-tRNA synthetase (EPRS), mRNA
M 004431	Homo sapiens EphA2 (EPHA2), mRNA Homo sapiens erythrocyte membrane protein band 7.2 (stomatin) (EPB72),
NM_004099	
T 5 00 1107	mRNA Homo sapiens erythrocyte membrane protein band 4.1 (elliptocytosis 1, RH-
NM_004437	
7.5.004425	linked) (EPB41), mRNA Homo sapiens endonuclease G (ENDOG), nuclear gene encoding mitochondrial
NM_004435	
ND 4 00 4424	Homo sapiens echinoderm microtubule-associated protein-like (EMAPL),
NM_004434	
NT 6 004422	mRNA Homo sapiens E74-like factor 3 (ets domain transcription factor, epithelial-
NM_004433	
> 7 C 004006	Homo sapiens eukaryotic translation initiation factor 4E binding protein 2
NM_004096	(
ND 6 004005	Homo sapiens eukaryotic translation initiation factor 4E binding protein 1
NM_004095	(FIE/ERPI) mRNA
>D f 004420	Homo sapiens early growth response 3 (EGR3), mRNA
NM_004430	Homo sapiens ephrin-B2 (EFNB2), mRNA
NM_004093	Homo sapiens ephrin-B1 (EFNB1), mRNA
NM_004429	Homo sapiens ephrin-A1 (EFNA1), mRNA  Homo sapiens ephrin-A1 (EFNA1), mRNA
NM_004428	Homo sapiens epinin-AT (EFTATT), indexing 2A (ITM2A), mRNA  Homo sapiens integral membrane protein 2A (ITM2A), mRNA
NM_004867	Try domeonicism (DPI DPII) (DSP), MKNA
NM_004415	Homo sapiens desmoplarin (DTI, DTID), Homo sapiens serine/threonine kinase 17a (apoptosis-inducing) (STK17A),
NM_004760	
	mRNA Homo sapiens dipeptidase 1 (renal) (DPEP1), mRNA
NM_004413	Homo sapiens dipeptidase 1 (Ichar) (BFES 1); terminal (DNTT), mRNA  Homo sapiens deoxynucleotidyltransferase, terminal (DNTT), mRNA
NM_004088	Homo sapiens deoxynucieotidylti aisterase, terratical (DNCI), mRNA  Homo sapiens DNA (cytosine-5-)-methyltransferase 2 (DNMT2), mRNA
NM_004412	Homo sapiens DNA (cytosine-5-)-inchryntalistete polypeptide 1 (DNCI1), mRNA Homo sapiens dynein, cytoplasmic, intermediate polypeptide 1 (DNCI1), mRNA
NM_004411	Homo sapiens dynein, cytopiasinic, intermediate posphorote (DMP1), mRNA  Homo sapiens dentin matrix acidic phosphoprotein (DMP1), mRNA
NM_004407	Homo sapiens denun matrix acture phosphoprotein (2007) Homo sapiens discs, large (Drosophila) homolog-associated protein 1
NM_004746	Homo sapiens discs, large (Diosophila) homolog assessment
	(DLGAP1), mRNA  Homo sapiens discs, large (Drosophila) homolog 5 (DLG5), mRNA
NM_004747	Homo sapiens discs, large (Drosophila) homolog 1 (DLG1), mRNA  Homo sapiens discs, large (Drosophila) homolog 1 (DLG1), mRNA
NM_004087	Homo sapiens discs, large (Drosophila) homolog 1 (2223)  Homo sapiens phorbolin (similar to apolipoprotein B mRNA editing protein)
NM_004900	Homo sapiens phorbolin (similar to aponipoprotein B internal programme and a saping property of the saping programme and a saping property of the saping propert
	(DJ742C19.2), mRNA  Homo sapiens neural precursor cell expressed, developmentally down-regulated
NM_004404	Homo sapiens neural precursor cell expressed, developments
	5 (NEDD5), mRNA  Homo sapiens DNA fragmentation factor, 40 kD, beta polypeptide (caspase-
NM_004402	Homo sapiens DNA fragmentation factor, 40 kB, 50th polypopular
	activated DNase) (DFFB), mRNA  Homo sapiens DNA fragmentation factor, 45 kD, alpha polypeptide (DFFA),
NM_004401	Homo sapiens DNA tragmentation factor, 45 kD, aspita postportation
	mRNA  DNA 1 in ducible transcript 3 (DDIT3), mRNA
NM_004083	Homo sapiens DNA-damage-inducible transcript 3 (DDIT3), mRNA  Homo sapiens DNA-damage-inducible transcript 3 (DDIT3), mRNA
NM_004734	Homo sapiens doublecortin and CaM kinase-like 1 (DCAMKL1), mRNA  Homo sapiens doublecortin and CaM kinase-like 1 (DCAMKL1), mRNA
NM_004394	Homo sapiens death-associated protein (DAP), mRNA  Homo sapiens death-associated protein (DAP), mRNA  (DAG1),
NM_004393	Homo sapiens death-associated protein (D/H), indeath Homo sapiens dystroglycan 1 (dystrophin-associated glycoprotein 1) (DAG1),
	mRNA i 16 G 1 to recriptional activation subunit 2
NM 004229	Homo sapiens cofactor required for Sp1 transcriptional activation, subunit 2
	(150kD) (CRSP2), mRNA
NM_004079	
NM 004390	Homo sapiens cathepsin H (CTSH), mRNA

	(compa) DATA
NM_004388_	Homo sapiens chitobiase, di-N-acetyl- (CTBS), mRNA
NM_004387	Homo sapiens cardiac-specific homeo box (CSX), mRNA
NM_004861	Homo sapiens cerebroside (3'-phosphoadenylylsulfate:galactosylceramide 3')
	sulfotransferase (CST), mRNA
NM_004078	Homo sapiens cysteine and glycine-rich protein 1 (CSRP1), mRNA
NM_004386	Homo sapiens chondroitin sulfate proteoglycan 3 (neurocan) (CSPG3), mRNA
NM_004385	Homo sapiens chondroitin sulfate proteoglycan 2 (versican) (CSPG2), mRNA
NM_004384	Homo sapiens casein kinase 1, gamma 3 (CSNK1G3), mRNA
NM 004383	Homo sapiens c-src tyrosine kinase (CSK), mRNA
NM 004075	Homo sapiens cryptochrome 1 (photolyase-like) (CRY1), mRNA
NM 004778	Homo sapiens G protein-coupled receptor 44 (GPR44), mRNA
NM 004750	Homo sapiens cytokine receptor-like factor 1 (CRLF1), mRNA
NM 004382	Homo sapiens corticotropin releasing hormone receptor 1 (CRHR1), mRNA
NM 004379	Homo sapiens cAMP responsive element binding protein 1 (CREB1), mRNA
NM 004377	Homo sapiens carnitine palmitoyltransferase I, muscle (CPT1B), mRNA
NM 004748	Homo sapiens cell cycle progression 8 protein (CPR8), mRNA
NM 004074	Homo sapiens cytochrome c oxidase subunit VIII (COX8), nuclear gene
11111_001071	encoding mitochondrial protein, mRNA
NM_004766	Homo sapiens coatomer protein complex, subunit beta 2 (beta prime) (COPB2),
	mRNA
NM 004645	Homo sapiens coilin (COIL), mRNA
NM 000614	Homo sapiens ciliary neurotrophic factor (CNTF), mRNA
NM 004368	Homo sapiens calponin 2 (CNN2), mRNA
NM 004072	Homo sapiens chemokine-like receptor 1 (CMKLR1), mRNA
NM 004071	Homo sapiens CDC-like kinase1 (CLK1), mRNA
NM 004362	Homo sapiens calmegin (CLGN), mRNA
NM 004070	Homo sapiens chloride channel Ka (CLCNKA), mRNA
NM 004804	Homo sapiens WD40 protein Ciao1 (CIAO1), mRNA
NM 004267	Homo sapiens carbohydrate (chondroitin 6/keratan) sulfotransferase 2 (CHST2),
1111_004207	mRNA
NM 004067	Homo sapiens chimerin (chimaerin) 2 (CHN2), mRNA
NM 004284	Homo sapiens chromodomain helicase DNA binding protein 1-like (CHD1L),
14141_00 120 1	mRNA
NM 004364	Homo sapiens CCAAT/enhancer binding protein (C/EBP), alpha (CEBPA),
11111_00 1501	mRNA
NM 004065	Homo sapiens cerebellar degeneration-related protein (34kD) (CDR1), mRNA
NM 004233	Homo sapiens CD83 antigen (activated B lymphocytes, immunoglobulin
14141_004255	superfamily) (CD83), mRNA
NM_004356	Homo sapiens CD81 antigen (target of antiproliferative antibody 1) (CD81),
14141_004550	mRNA
NM 004357	Homo sapiens CD151 antigen (CD151), mRNA
NM 004350	Homo sapiens cunt-related transcription factor 3 (RUNX3), mRNA
NM 004349	Homo sapiens core-binding factor, runt domain, alpha subunit 2; translocated to,
14141 004243	1; cyclin D-related (CBFA2T1), mRNA
NM 004345	Homo sapiens cathelicidin antimicrobial peptide (CAMP), mRNA
NM 000722	Homo sapiens catheneidin antimetobiai peptide (CAMP), inkiva  Homo sapiens calcium channel, voltage-dependent, alpha 2/delta subunit 1
14141_000/22	(CACNA2D1), mRNA
NM 004334	Homo sapiens bone marrow stromal cell antigen 1 (BST1), mRNA
	Homo sapiens small inducible cytokine subfamily B (Cys-X-Cys), member 14
NM_004887	(BRAK) (SCYB14), mRNA
NIM 004222	Homo sapiens v-raf murine sarcoma viral oncogene homolog B1 (BRAF),
NM_004333	mRNA
L	IIIVIVA

	Homo sapiens bone morphogenetic protein receptor, type IA (BMPR1A), mRNA
	Homo sapiens ATP-binding cassette, sub-family G (WHITE), member 2
NM_004827	(ABCG2), mRNA
NM 004326	Homo sapiens B-cell CLL/lymphoma 9 (BCL9), mRNA
NM 004765	Homo saniens B-cell CLL/lymphoma 7C (BCL/C), mRNA
NM 004324	Home saniens BCI 2-associated X protein (BAX), mRNA
NM 004656	Homo sapiens BRCA1 associated protein-1 (ubiquitin carboxy-terminal
14171_00 1050	hydrolase) (BAP1), mRNA
NM 004048	Homo sapiens beta-2-microglobulin (B2M), mRNA
NM 004655	Homo saniens axin 2 (conductin, axil) (AXIN2), mRNA
NM 004321	Homo saniens ayonal transport of synaptic vesicles (ATSV), mRNA
NM_004888	Homo sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump),
NM_004047	Homo sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump)
NM_004046	21kD (ATP6F), mRNA  Homo sapiens ATP synthase, H+ transporting, mitochondrial F1 complex, alpha subunit, isoform 1, cardiac muscle (ATP5A1), mRNA
NM_001683	Homo sapiens ATPase, Ca++ transporting, plasma membrane 2 (ATP2B2),
ND 4 004214	mRNA Homo sapiens ADP-ribosyltransferase 1 (ART1), mRNA
NM 004314	Homo sapiens arrestin, beta 2 (ARRB2), mRNA
NM_004313	Homo sapiens arrestin 3, retinal (X-arrestin) (ARR3), mRNA
NM_004312	Homo sapiens ADP-ribosylation factor-like 3 (ARL3), mRNA
NM_004311	Homo sapiens ras homolog gene family, member I (ARHI), mRNA
NM_004675	Homo sapiens ras homolog gene family, member H (ARHH), mRNA  Homo sapiens ras homolog gene family, member H (ARHH), mRNA
NM_004310	Homo sapiens Rho GDP dissociation inhibitor (GDI) alpha (ARHGDIA), mRNA
NM_004309	Homo sapiens Rho GTP dissociation infinition (GBs) diplies (Fig. 1). Homo sapiens Rho GTPase activating protein 1 (ARHGAP1), mRNA
NM_004308	Homo sapiens ras homolog gene family, member B (ARHB), mRNA
NM_004040	Homo sapiens ras nomolog gene laminy, member B (radib), member B (
NM_004290	Homo sapiens ring finger protein 14 (RNF14), mRNA
NM_004797	Homo sapiens adipose most abundant gene transcript 1 (APM1), mRNA
NM_004039	Homo sapiens annexin A2 (ANXA2), mRNA
NM_004306	Homo sapiens annexin A13 (ANXA13), mRNA
NM_004038	Homo sapiens amylase, alpha 1A; salivary (AMY1A), mRNA
NM_004305	Homo sapiens bridging integrator 1 (BIN1), mRNA
NM_004857	Homo sapiens A kinase (PRKA) anchor protein 5 (AKAP5), mRNA
NM_004833	Homo sapiens absent in melanoma 2 (AIM2), mRNA
NM_004208	Homo sapiens programmed cell death 8 (apoptosis-inducing factor) (PDCD8), mRNA
NM 002199	Homo sapiens interferon regulatory factor 2 (IRF2), mRNA
NM 001569	Homo saniens interleukin-1 receptor-associated kinase I (IRAK1), mRNA
NM 001567	Home saniens inesited polyphosphate phosphatase-like I (INPPLI), mRNA
NM 002194	Homo sapiens inositol polyphosphate-1-phosphatase (INPPI), mRNA
NM 002111	Homo seniens huntingtin (Huntington disease) (HD), mRNA
NM 000165	Homo saniens gan junction protein, alpha 1, 43kD (connexin 43) (GJA1), mRNA
NM_001999	Homo sapiens fibrillin 2 (congenital contractural arachnodactyly) (FBN2), mRNA
NM 001937	Homo saniens dermatopontin (DPT), mRNA
NM_001381	Homo sapiens docking protein 1, 62kD (downstream of tyrosine kinase 1) (DOK1), mRNA
NM_000729	Homo sapiens cholecystokinin (CCK), mRNA
NM_000486	Homo saniens aquanorin 2 (collecting duct) (AQP2), mRNA
NM 001520	
14141_001320	Trous and

<del></del>	2001 D.) (CTE2C1) DNIA
77.6 000007	220kD) (GTF3C1), mRNA
NM_002097	Homo sapiens general transcription factor IIIA (GTF3A), mRNA
NM_003205	Homo sapiens transcription factor 12 (HTF4, helix-loop-helix transcription
	factors 4) (TCF12), mRNA
NM_000440	Homo sapiens phosphodiesterase 6A, cGMP-specific, rod, alpha (PDE6A),
- <del></del>	mRNA
NM_000806	Homo sapiens gamma-aminobutyric acid (GABA) A receptor, alpha 1
	(GABRA1), mRNA
NM_001809	Homo sapiens centromere protein A (17kD) (CENPA), mRNA
NM_000439	Homo sapiens proprotein convertase subtilisin/kexin type 1 (PCSK1), mRNA
NM_002529	Homo sapiens neurotrophic tyrosine kinase, receptor, type 1 (NTRK1), mRNA
NM_003417	Homo sapiens zinc finger protein 264 (ZNF264), mRNA
NM_000395	Homo sapiens colony stimulating factor 2 receptor, beta, low-affinity
	(granulocyte-macrophage) (CSF2RB), mRNA
NM_000065	Homo sapiens complement component 6 (C6), mRNA
NM_000252	Homo sapiens myotubular myopathy 1 (MTM1), mRNA
NM_000229	Homo sapiens lecithin-cholesterol acyltransferase (LCAT), nuclear gene
	encoding mitochondrial protein, mRNA
NM_000224	Homo sapiens keratin 18 (KRT18), mRNA
NM_000211	Homo sapiens integrin, beta 2 (antigen CD18 (p95), lymphocyte function-
	associated antigen 1; macrophage antigen 1 (mac-1) beta subunit) (ITGB2),
	mRNA
NM_000208	Homo sapiens insulin receptor (INSR), mRNA
NM 000206	Homo sapiens interleukin 2 receptor, gamma (severe combined
-	immunodeficiency) (IL2RG), mRNA
NM 000416	Homo sapiens interferon gamma receptor 1 (IFNGR1), mRNA
NM 000201	Homo sapiens intercellular adhesion molecule 1 (CD54), human rhinovirus
-	receptor (ICAM1), mRNA
NM_000350	Homo sapiens ATP-binding cassette, sub-family A (ABC1), member 4
	(ABCA4), mRNA
NM 000110	Homo sapiens dihydropyrimidine dehydrogenase (DPYD), mRNA
NM_000375	Homo sapiens uroporphyrinogen III synthase (congenital erythropoietic
_	porphyria) (UROS), mRNA
NM 000459	Homo sapiens TEK tyrosine kinase, endothelial (venous malformations, multiple
_	cutaneous and mucosal) (TEK), mRNA
NM 001053	Homo sapiens somatostatin receptor 5 (SSTR5), mRNA
NM 001052	Homo sapiens somatostatin receptor 4 (SSTR4), mRNA
NM 001051	Homo sapiens somatostatin receptor 3 (SSTR3), mRNA
NM_001050	Homo sapiens somatostatin receptor 2 (SSTR2), mRNA
NM 001049	Homo sapiens somatostatin receptor 1 (SSTR1), mRNA
NM 000348	Homo sapiens steroid-5-alpha-reductase, alpha polypeptide 2 (3-oxo-5 alpha-
	steroid delta 4-dehydrogenase alpha 2) (SRD5A2), mRNA
NM 000340	Homo sapiens solute carrier family 2 (facilitated glucose transporter), member 2
	(SLC2A2), mRNA
NM_000338	Homo sapiens solute carrier family 12 (sodium/potassium/chloride transporters),
	member 1 (SLC12A1), mRNA
NM 000231	Homo sapiens sarcoglycan, gamma (35kD dystrophin-associated glycoprotein)
1111_000251	(SGCG), mRNA
NM 001034	Homo sapiens ribonucleotide reductase M2 polypeptide (RRM2), mRNA
NM 000448	Homo sapiens recombination activating gene 1 (RAG1), mRNA
NM 000303	Homo sapiens phosphomannomutase 2 (PMM2), mRNA
NM 000303	Homo sapiens procollagen-lysine, 2-oxoglutarate 5-dioxygenase (lysine
14147 000205	110mo sapiens proconagen-rysme, 2-oxogranate 3-dioxygenase (rysme

	hydroxylase, Ehlers-Danlos syndrome type VI) (PLOD), mRNA
	ry arrangemental Coentyme A Caluary and a result in
VM_000282	nuclear gene encoding mitochondrial protein, mRNA
	Homo sapiens 6-pyruvoyl-tetrahydropterin synthase/dimerization cofactor of
VM_000281	Homo sapiens 6-pyruvoyi-tettariydi operatory (PCBD), mRNA hepatocyte nuclear factor 1 alpha (TCF1) (PCBD), mRNA
NM_000277	Homo sapiens phenylalanine hydroxylase (1711), interest Homo sapiens 3-oxoacid CoA transferase (OXCT), nuclear gene encoding
NM_000436	Homo sapiens 3-oxoacid CoA transicrase (GIZGZ),
	mitochondrial protein, mRNA  Homo sapiens ornithine aminotransferase (gyrate atrophy) (OAT), nuclear gene
NM_000274	Homo sapiens ornithine aminouansiciase (gyrate and r
	encoding mitochondrial protein, mRNA  Homo sapiens ocular albinism 1 (Nettleship-Falls) (OA1), mRNA  ORDHRI) mRNA
NM_000273	Homo sapiens ocular albinism i (Neurosino 1 dinyenile) (NPHP1), mRNA
NM 000272	Homo sapiens octular aloinism i (Nemory), mRNA  Homo sapiens nephronophthisis 1 (juvenile) (NPHP1), mRNA
NM 000271	Homo sapiens Niemann-Pick disease, type C1 (NPC1), mRNA  Homo sapiens non-metastatic cells 1, protein (NM23A) expressed in (NME1),
NM_000269	Homo sapiens non-metastatic cells 1, protein (1VIVI2311) outprotein
- ·- · · <del>-</del>	
NM 000268	mRNA Homo sapiens neurofibromin 2 (bilateral acoustic neuroma) (NF2), mRNA  Glacometeris, you Recklinghausen disease,
NM_000267	Homo sapiens neurofibromin I (neurofibromatosis, von Recenting
	1 - 1: \ (A)T(1) ma(1) (A) (A)
NM 000434	in a giolidose 1 (lycosomal statidase) (NEO 1), inicial
NM 000266	
NM_000265	Homo saniens neutrophil cytosolic factor 1 (4/kD), chromo granation
14141_000200	1 1) /NCETY mR NA
NM 000262	
NM_000261	Homo sapiens N-acetylgalactosaminidase, alpha (1776-17).  Homo sapiens myocilin, trabecular meshwork inducible glucocorticoid response
14141_000201	
NM_000258	(MYOC), mRNA  Homo sapiens myosin, light polypeptide 3, alkali; ventricular, skeletal, slow
14141_000250	(MYL3), mRNA
NM_000432	(MYL3), mRNA Homo sapiens myosin, light polypeptide 2, regulatory, cardiac, slow (MYL2),
NW_000432	
NM_000257	mRNA Homo sapiens myosin, heavy polypeptide 7, cardiac muscle, beta (MYH7),
14M_000257	
NM 000431	- lengte kingse (meyalonic aciduria) (MVN), likiva
NM 000451	Homo saniens methylmalonyl Coenzyme A mutase (WOT), haddan g
NM_000255	
27.6 000254	encoding mitochondrial protein, mRNA  Homo sapiens 5-methyltetrahydrofolate-homocysteine methyltransferase (MTR)
NM_000254	mPNA
37.5 000252	mRNA Homo sapiens microsomal triglyceride transfer protein (large polypeptide, 88kD)
NM_000253	
27.4 000250	(MTP), mRNA  Homo sapiens myeloperoxidase (MPO), nuclear gene encoding mitochondrial
NM_000250	
37.5 000248	
NM_000248	TY MIC class I polypentide-related sequence A (MC12);
NM_000247	
NM 000246	
NM_000245	DATA
	mRNA Homo sapiens multiple endocrine neoplasia I (MEN1), mRNA
NM_000244	
NM_000243	
NM_000242	
	(MBL2), mRNA  Homo sapiens methionine adenosyltransferase I, alpha (MAT1A), mRNA  Office of the control of the
NM_000429	
NM_00024	Homo sapiens monoamine oxidase A (MI 1011), Madical games
<u> </u>	mitochondrial protein, mRNA

NM_000428	Homo sapiens latent transforming growth factor beta binding protein 2 (LTBP2), mRNA
NM_000238	Homo sapiens potassium voltage-gated channel, subfamily H (eag-related), member 2 (KCNH2), mRNA
NM 000237	Homo sapiens lipoprotein lipase (LPL), mRNA
NM 000427	Homo sapiens loricrin (LOR), mRNA
NM 000236	Homo sapiens lipase, hepatic (LIPC), mRNA
NM 000235	Homo sapiens lipase A, lysosomal acid, cholesterol esterase (Wolman disease)
14141_000255	(LIPA), mRNA
NM 000234	Homo sapiens ligase I, DNA, ATP-dependent (LIG1), mRNA
NM_000233	Homo sapiens luteinizing hormone/choriogonadotropin receptor (LHCGR), mRNA
NM 000228	Homo sapiens laminin, beta 3 (nicein (125kD), kalinin (140kD), BM600
- system	(125kD)) (LAMB3), mRNA
NM_000426	Homo sapiens laminin, alpha 2 (merosin, congenital muscular dystrophy) (LAMA2), mRNA
NM_000226	Homo sapiens keratin 9 (epidermolytic palmoplantar keratoderma) (KRT9),
	mRNA
NM_000422	Homo sapiens keratin 17 (KRT17), mRNA
NM_000223	Homo sapiens keratin 12 (Meesmann corneal dystrophy) (KRT12), mRNA
NM_000421	Homo sapiens keratin 10 (epidermolytic hyperkeratosis; keratosis palmaris et plantaris) (KRT10), mRNA
NM_000222	Homo sapiens v-kit Hardy-Zuckerman 4 feline sarcoma viral oncogene homolog (KIT), mRNA
NM_000218	Homo sapiens potassium voltage-gated channel, KQT-like subfamily, member 1 (KCNQ1), mRNA
NM_000219	Homo sapiens potassium voltage-gated channel, Isk-related family, member 1 (KCNE1), mRNA
NM_000217	Homo sapiens potassium voltage-gated channel, shaker-related subfamily, member 1 (episodic ataxia with myokymia) (KCNA1), mRNA
NM 000216	Homo sapiens Kallmann syndrome 1 sequence (KAL1), mRNA
NM_000215	Homo sapiens Janus kinase 3 (a protein tyrosine kinase, leukocyte) (JAK3), mRNA
NM_000212	Homo sapiens integrin, beta 3 (platelet glycoprotein IIIa, antigen CD61) (ITGB3), mRNA
NM_000209	Homo sapiens insulin promoter factor 1, homeodomain transcription factor (IPF1), mRNA
NM 000207	Homo sapiens insulin (INS), mRNA
NM 000418	Homo sapiens interleukin 4 receptor (IL4R), mRNA
NM 000417	Homo sapiens interleukin 2 receptor, alpha (IL2RA), mRNA
NM 001551	Homo sapiens immunoglobulin (CD79A) binding protein 1 (IGBP1), mRNA
NM 000203	Homo sapiens iduronidase, alpha-L- (IDUA), mRNA
NM 000415	Homo sapiens islet amyloid polypeptide (IAPP), mRNA
NM 000200	Homo sapiens histatin 3 (HTN3), mRNA
NM 001538	Homo sapiens heat shock transcription factor 4 (HSF4), mRNA
NM_000859	Homo sapiens 3-hydroxy-3-methylglutaryl-Coenzyme A reductase (HMGCR), mRNA
NM 001527	Homo sapiens histone deacetylase 2 (HDAC2), mRNA
NM 001525	Homo sapiens hypocretin (orexin) receptor 1 (HCRTR1), mRNA
NM 001524	Homo sapiens hypocretin (orexin) neuropeptide precursor (HCRT), mRNA
NM 001510	Homo sapiens glutamate receptor, ionotropic, delta 2 (GRID2), mRNA
NM 000829	Homo sapiens glutamate receptor, ionotrophic, AMPA 4 (GRIA4), mRNA
TATAT 000973	1 Monto sapione giutamate receptor, fononopino, 1244 114 (Ozari), matri

	•
NM_001496	Homo sapiens GDNF family receptor alpha 3 (GFRA3), mRNA
NIM 001486	Homo saniens glucokinase (hexokinase 4) regulatory protein (GCKR), IIINNA
NTM ( 0000000	Homo soniens growth arrest-specific 6 (GAS6), mRNA
NM 000155	Home seniens galactose-1-phosphate uridylyltransterase (GALI), mRNA
NM 000153	Home seniers galactosylceramidase (Krabbe disease) (GALC), mKNA
NM_000816	Homo sapiens gamma-aminobutyric acid (GABA) A receptor, gamma 2
14141_000010	(CADDC2) mDNA
NM_000815	Homo sapiens gamma-aminobutyric acid (GABA) A receptor, delta (GABRD),
14141_000912	mDNA
ND 4 000911	Homo sapiens gamma-aminobutyric acid (GABA) A receptor, alpha 6
NM_000811	(CARPA6) mRNA
NR 4 000000	Homo sapiens gamma-aminobutyric acid (GABA) A receptor, alpha 4
NM_000809	(CARRAI) mRNA
ND 4 000000	Homo sapiens gamma-aminobutyric acid (GABA) A receptor, alpha 3
NM_000808	(CADDA3) mRNA
>7 f 000007	Homo sapiens gamma-aminobutyric acid (GABA) A receptor, alpha 2
NM_000807	(CADDA2) mDNA
77.6 000151	Homo sapiens glucose-6-phosphatase, catalytic (glycogen storage disease type I,
NM_000151	von Gierke disease) (G6PC), mRNA
27.5.001450	Homo sapiens forkhead box F2 (FOXF2), mRNA
NM_001452	Homo sapiens fibrillin 1 (Marfan syndrome) (FBN1), mRNA
NM_000138	Homo sapiens Fanconi anemia, complementation group C (FANCC), mRNA  Homo sapiens Fanconi anemia, complementation group C (FANCC), mRNA
NM_000136	Homo sapiens fatty acid binding protein 6, ileal (gastrotropin) (FABP6), mRNA
NM_001445	Homo sapiens fatty acid binding protein 4, adipocyte (FABP4), mRNA  Homo sapiens fatty acid binding protein 4, adipocyte (FABP4), mRNA
NM_001442	Homo sapiens fatty acid binding protein 4, adipocyte (1715), india
NM_001443	Homo sapiens fatty acid binding protein 1, liver (FABP1), mRNA
NM_001441	Homo sapiens fatty acid amide hydrolase (FAAH), mRNA
NM_000401	Homo sapiens exostoses (multiple) 2 (EXT2), mRNA
NM_000127	Homo sapiens exostoses (multiple) 1 (EXT1), mRNA
NM_001433	Homo sapiens ER to nucleus signalling 1 (ERN1), mRNA
NM_000122	Homo sapiens excision repair cross-complementing rodent repair deficiency,
	complementation group 3 (xeroderma pigmentosum group B complementing)
	(ERCC3), mRNA
NM_000121	Homo sapiens erythropoietin receptor (EPOR), mRNA
NM_000120	Homo sapiens epoxide hydrolase 1, microsomal (xenobiotic) (EPHX1), mRNA
NM 000119	Homo sapiens erythrocyte membrane protein band 4.2 (EPB42), mRNA
NM 001429	Llome senions E1 A hinding protein p300 (EP300), mKNA
NM 000118	Homo saniens endoglin (Osler-Rendu-Weber syndrome 1) (ENG), IIIKNA
NM_000117	Homo saniens emerin (Emery-Dreifuss muscular dystrophy) (EVID), IIINIA
NM 001422	Homo sapiens E74-like factor 5 (ets domain franscription factor) (ELF3), filk NA
NM 000114	II-ma ranians and othelin 3 (FDN3) mRNA
NM_001393	Homo sapiens extracellular matrix protein 2, female organ and adipocyte specific
_	(FCM2) mRNA
NM_000112	Homo sapiens solute carrier family 26 (sulfate transporter), member 2
	(SLC26A2), mRNA
NM 001382	Homo saniens dolichyl-phosphate (UDP-N-acetylglucosamine) N-
	acetylglucosaminephosphotransferase 1 (GlcNAc-1-P transferase) (DPAGT1),
	mRNA
NM 001365	Homo sapiens discs, large (Drosophila) homolog 4 (DLG4), mRNA
NM_000792	Homo sapiens deiodinase, jodothyronine, type I (DIOI), mRNA
NM 001358	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 15 (DDX15),
1444-001550	mPNA
1	Homo sapiens damage-specific DNA binding protein 2 (48kD) (DDB2), mRNA

NM 00101	er 2
NM_001334   Homo sapiens cubilin (intrinsic factor-cobalamin receptor) (CUBN), mR NM_001334   Homo sapiens cathepsin O (CTSO), mRNA	er 2
NM 001334	er 2
NM 001334	er 2
NM_000554Homo sapiens cone-rod homeobox (CRX), mRNANM_000096Homo sapiens ceruloplasmin (ferroxidase) (CP), mRNANM_000095Homo sapiens cartilage oligomeric matrix protein (pseudoachondroplasia epiphyseal dysplasia 1, multiple) (COMP), mRNANM_000392Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member (ABCC2), mRNANM_000085Homo sapiens chloride channel Kb (CLCNKB), mRNANM_000084Homo sapiens chloride channel 5 (nephrolithiasis 2, X-linked, Dent disea (CLCN5), mRNANM_001279Homo sapiens cell death-inducing DFFA-like effector a (CIDEA), mRNNM_000080Homo sapiens cholinergic receptor, nicotinic, epsilon polypeptide (CHR mRNANM_000751Homo sapiens cholinergic receptor, nicotinic, delta polypeptide (CHRNI mRNANM_000747Homo sapiens cholinergic receptor, nicotinic, beta polypeptide 1 (muscle (CHRNA1), mRNANM_00079Homo sapiens cholimergic receptor, nicotinic, alpha polypeptide 1 (muscle (CHRNA1), mRNANM_001271Homo sapiens chromodomain helicase DNA binding protein 4 (CHD4), NM 001270NM_001270Homo sapiens chromodomain helicase DNA binding protein 1 (CHD1), NM 000078NM_00078Homo sapiens cholesteryl ester transfer protein, plasma (CETP), mRNANM_000078Homo sapiens cyclin-dependent kinase inhibitor 1C (p57, Kip2) (CDKn mRNANM_001251Homo sapiens CD68 antigen (CD68), mRNANM_000074Homo sapiens CD68 antigen (CD68), mRNANM_000073Homo sapiens CD3G antigen, gamma polypeptide (TiT3 complex) (CD mRNA	er 2
NM_000554         Homo sapiens cone-rod homeobox (CRX), mRNA           NM_000096         Homo sapiens ceruloplasmin (ferroxidase) (CP), mRNA           NM_000095         Homo sapiens cartilage oligomeric matrix protein (pseudoachondroplasia epiphyseal dysplasia 1, multiple) (COMP), mRNA           NM_000392         Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), memb (ABCC2), mRNA           NM_000085         Homo sapiens chloride channel Kb (CLCNKB), mRNA           NM_000084         Homo sapiens chloride channel 5 (nephrolithiasis 2, X-linked, Dent disea (CLCN5), mRNA           NM_001279         Homo sapiens cell death-inducing DFFA-like effector a (CIDEA), mRN           NM_000080         Homo sapiens cholinergic receptor, nicotinic, epsilon polypeptide (CHR mRNA           NM_000751         Homo sapiens cholinergic receptor, nicotinic, delta polypeptide 1 (muscle (CHRNB1), mRNA           NM_000794         Homo sapiens cholinergic receptor, nicotinic, beta polypeptide 1 (muscle (CHRNA1), mRNA           NM_001271         Homo sapiens chromodomain helicase DNA binding protein 4 (CHD4), NM 001271           NM_001270         Homo sapiens chromodomain helicase DNA binding protein 1 (CHD1), NM 000078           NM_001271         Homo sapiens chromodomain helicase DNA binding protein 1 (CHD1), NM 000078           NM_001251         Homo sapiens cyclin-dependent kinase inhibitor 1C (p57, Kip2) (CDKn mRNA)           NM_001251         Homo sapiens CD68 antigen (CD68), mRNA	er 2
NM_000096   Homo sapiens ceruloplasmin (ferroxidase) (CP), mRNA	er 2
NM_000395	er 2
epiphyseal dysplasia 1, multiple) (COMP), mRNA  NM_000392 Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), members (ABCC2), mRNA  NM_000085 Homo sapiens chloride channel Kb (CLCNKB), mRNA  NM_000084 Homo sapiens chloride channel 5 (nephrolithiasis 2, X-linked, Dent disertion (CLCN5), mRNA  NM_001279 Homo sapiens cell death-inducing DFFA-like effector a (CIDEA), mRNA  NM_000080 Homo sapiens cholinergic receptor, nicotinic, epsilon polypeptide (CHRNImRNA  NM_000751 Homo sapiens cholinergic receptor, nicotinic, delta polypeptide (CHRNImRNA  NM_000747 Homo sapiens cholinergic receptor, nicotinic, beta polypeptide 1 (muscle (CHRNB1), mRNA  NM_000079 Homo sapiens cholinergic receptor, nicotinic, alpha polypeptide 1 (muscle (CHRNA1), mRNA  NM_001273 Homo sapiens chromodomain helicase DNA binding protein 4 (CHD4), NM_001271 Homo sapiens chromodomain helicase DNA binding protein 2 (CHD2), NM 001270 Homo sapiens chromodomain helicase DNA binding protein 1 (CHD1), NM 000078 Homo sapiens cholesteryl ester transfer protein, plasma (CETP), mRNA  NM_001075 Homo sapiens cyclin-dependent kinase inhibitor 1C (p57, Kip2) (CDKN mRNA  NM_001251 Homo sapiens CD68 antigen (CD68), mRNA  NM_000074 Homo sapiens CD68 antigen (CD68), mRNA  NM_000075 Homo sapiens CD68 antigen (CD68), mRNA  NM_000076 Homo sapiens CD68 antigen (CD68), mRNA  NM_000077 Homo sapiens CD68 antigen (CD68), mRNA  NM_000078 Homo sapiens CD68 antigen (CD68), mRNA  NM_000079 Homo sapiens CD68 antigen (CD68), mRNA  NM_000079 Homo sapiens CD68 antigen (CD68), mRNA	er 2
NM_000392 Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member (ABCC2), mRNA  NM_000085 Homo sapiens chloride channel Kb (CLCNKB), mRNA  NM_000084 Homo sapiens chloride channel 5 (nephrolithiasis 2, X-linked, Dent diser (CLCN5), mRNA  NM_001279 Homo sapiens cell death-inducing DFFA-like effector a (CIDEA), mRNA  NM_000080 Homo sapiens cholinergic receptor, nicotinic, epsilon polypeptide (CHR mRNA  NM_000751 Homo sapiens cholinergic receptor, nicotinic, delta polypeptide (CHRNI mRNA  NM_000747 Homo sapiens cholinergic receptor, nicotinic, beta polypeptide 1 (muscle (CHRNB1), mRNA  NM_000079 Homo sapiens cholinergic receptor, nicotinic, alpha polypeptide 1 (muscle (CHRNA1), mRNA  NM_001273 Homo sapiens chromodomain helicase DNA binding protein 4 (CHD4), NM_001271 Homo sapiens chromodomain helicase DNA binding protein 2 (CHD2), NM_001270 Homo sapiens chromodomain helicase DNA binding protein 1 (CHD1), NM_000078 Homo sapiens cholesteryl ester transfer protein, plasma (CETP), mRNA  NM_0010251 Homo sapiens cyclin-dependent kinase inhibitor 1C (p57, Kip2) (CDKN mRNA  NM_001251 Homo sapiens CD68 antigen (CD68), mRNA  NM_00100073 Homo sapiens CD68 antigen (CD68), mRNA  NM_000073 Homo sapiens CD68 antigen (CD68), mRNA  NM_000073 Homo sapiens CD68 antigen, gamma polypeptide (TiT3 complex) (CD mRNA	ase)
NM_000085Homo sapiens chloride channel Kb (CLCNKB), mRNANM_000084Homo sapiens chloride channel 5 (nephrolithiasis 2, X-linked, Dent disea (CLCN5), mRNANM_001279Homo sapiens cell death-inducing DFFA-like effector a (CIDEA), mRNNM_000080Homo sapiens cholinergic receptor, nicotinic, epsilon polypeptide (CHR mRNANM_000751Homo sapiens cholinergic receptor, nicotinic, delta polypeptide (CHRNI mRNANM_000747Homo sapiens cholinergic receptor, nicotinic, beta polypeptide 1 (muscle (CHRNB1), mRNANM_000079Homo sapiens cholinergic receptor, nicotinic, alpha polypeptide 1 (muscle (CHRNA1), mRNANM_01273Homo sapiens chromodomain helicase DNA binding protein 4 (CHD4), NM 001271NM_001270Homo sapiens chromodomain helicase DNA binding protein 2 (CHD2), NM 001270NM_001270Homo sapiens cholesteryl ester transfer protein, plasma (CETP), mRNANM_000078Homo sapiens cyclin-dependent kinase inhibitor 1C (p57, Kip2) (CDKN mRNANM_001251Homo sapiens cyclin-dependent kinase 3 (CDK3), mRNANM_001251Homo sapiens CD68 antigen (CD68), mRNANM_000074Homo sapiens tumor necrosis factor (ligand) superfamily, member 5 (hy syndrome) (TNFSF5), mRNANM_000073Homo sapiens CD3G antigen, gamma polypeptide (TiT3 complex) (CD mRNA	
NM_000084 Homo sapiens chloride channel 5 (nephrolithiasis 2, X-linked, Dent disea (CLCN5), mRNA  NM_001279 Homo sapiens cell death-inducing DFFA-like effector a (CIDEA), mRN NM_000080 Homo sapiens cholinergic receptor, nicotinic, epsilon polypeptide (CHR mRNA  NM_000751 Homo sapiens cholinergic receptor, nicotinic, delta polypeptide (CHRNI mRNA  NM_000747 Homo sapiens cholinergic receptor, nicotinic, beta polypeptide 1 (muscle (CHRNB1), mRNA  NM_000079 Homo sapiens cholinergic receptor, nicotinic, alpha polypeptide 1 (muscle (CHRNA1), mRNA  NM_001273 Homo sapiens chromodomain helicase DNA binding protein 4 (CHD4), NM_001271 Homo sapiens chromodomain helicase DNA binding protein 2 (CHD2), NM_001270 Homo sapiens chromodomain helicase DNA binding protein 1 (CHD1), NM_000078 Homo sapiens cholesteryl ester transfer protein, plasma (CETP), mRNA  NM_000076 Homo sapiens cyclin-dependent kinase inhibitor 1C (p57, Kip2) (CDKN mRNA  NM_001251 Homo sapiens CD68 antigen (CD68), mRNA  NM_000074 Homo sapiens tumor necrosis factor (ligand) superfamily, member 5 (hy syndrome) (TNFSF5), mRNA  NM_000073 Homo sapiens CD3G antigen, gamma polypeptide (TiT3 complex) (CD mRNA)	
CLCN5), mRNA	
NM_001279 Homo sapiens cell death-inducing DFFA-like effector a (CIDEA), mRN NM_000080 Homo sapiens cholinergic receptor, nicotinic, epsilon polypeptide (CHR mRNA  NM_000751 Homo sapiens cholinergic receptor, nicotinic, delta polypeptide (CHRNI mRNA  NM_000747 Homo sapiens cholinergic receptor, nicotinic, beta polypeptide 1 (muscle (CHRNB1), mRNA  NM_000079 Homo sapiens cholinergic receptor, nicotinic, alpha polypeptide 1 (muscle (CHRNA1), mRNA  NM_001273 Homo sapiens chromodomain helicase DNA binding protein 4 (CHD4), NM_001271 Homo sapiens chromodomain helicase DNA binding protein 2 (CHD2), NM_001270 Homo sapiens chromodomain helicase DNA binding protein 1 (CHD1), NM_000078 Homo sapiens cholesteryl ester transfer protein, plasma (CETP), mRNA  NM_000076 Homo sapiens cyclin-dependent kinase inhibitor 1C (p57, Kip2) (CDKN mRNA  NM_001251 Homo sapiens cyclin-dependent kinase 3 (CDK3), mRNA  NM_000074 Homo sapiens tumor necrosis factor (ligand) superfamily, member 5 (hy syndrome) (TNFSF5), mRNA  NM_000073 Homo sapiens CD3G antigen, gamma polypeptide (TiT3 complex) (CD mRNA	A
NM_000080 Homo sapiens cholinergic receptor, nicotinic, epsilon polypeptide (CHR) mRNA  NM_000751 Homo sapiens cholinergic receptor, nicotinic, delta polypeptide (CHRNI mRNA  NM_000747 Homo sapiens cholinergic receptor, nicotinic, beta polypeptide 1 (muscle (CHRNB1), mRNA  NM_000079 Homo sapiens cholinergic receptor, nicotinic, alpha polypeptide 1 (muscle (CHRNA1), mRNA  NM_001273 Homo sapiens chromodomain helicase DNA binding protein 4 (CHD4), NM_001271 Homo sapiens chromodomain helicase DNA binding protein 2 (CHD2), NM_001270 Homo sapiens chromodomain helicase DNA binding protein 1 (CHD1), NM_000078 Homo sapiens cholesteryl ester transfer protein, plasma (CETP), mRNA  NM_000076 Homo sapiens cyclin-dependent kinase inhibitor 1C (p57, Kip2) (CDKN mRNA  NM_001251 Homo sapiens CD68 antigen (CD68), mRNA  NM_000074 Homo sapiens tumor necrosis factor (ligand) superfamily, member 5 (hy syndrome) (TNFSF5), mRNA  NM_000073 Homo sapiens CD3G antigen, gamma polypeptide (TiT3 complex) (CD mRNA	
mRNA  NM_000751 Homo sapiens cholinergic receptor, nicotinic, delta polypeptide (CHRNI mRNA  NM_000747 Homo sapiens cholinergic receptor, nicotinic, beta polypeptide 1 (muscle (CHRNB1), mRNA  NM_000079 Homo sapiens cholinergic receptor, nicotinic, alpha polypeptide 1 (muscle (CHRNA1), mRNA  NM_001273 Homo sapiens chromodomain helicase DNA binding protein 4 (CHD4), NM_001271 Homo sapiens chromodomain helicase DNA binding protein 2 (CHD2), NM_001270 Homo sapiens chromodomain helicase DNA binding protein 1 (CHD1), NM_000078 Homo sapiens cholesteryl ester transfer protein, plasma (CETP), mRNA  NM_000076 Homo sapiens cyclin-dependent kinase inhibitor 1C (p57, Kip2) (CDKN mRNA  NM_001251 Homo sapiens CD68 antigen (CD68), mRNA  NM_000074 Homo sapiens tumor necrosis factor (ligand) superfamily, member 5 (hy syndrome) (TNFSF5), mRNA  NM_000073 Homo sapiens CD3G antigen, gamma polypeptide (TiT3 complex) (CD mRNA)	
NM_000751 Homo sapiens cholinergic receptor, nicotinic, delta polypeptide (CHRNI mRNA  NM_000747 Homo sapiens cholinergic receptor, nicotinic, beta polypeptide 1 (muscle (CHRNB1), mRNA  NM_000079 Homo sapiens cholinergic receptor, nicotinic, alpha polypeptide 1 (muscle (CHRNA1), mRNA  NM_001273 Homo sapiens chromodomain helicase DNA binding protein 4 (CHD4), NM_001271 Homo sapiens chromodomain helicase DNA binding protein 2 (CHD2), NM_001270 Homo sapiens chromodomain helicase DNA binding protein 1 (CHD1), NM_000078 Homo sapiens cholesteryl ester transfer protein, plasma (CETP), mRNA  NM_000076 Homo sapiens cyclin-dependent kinase inhibitor 1C (p57, Kip2) (CDKN mRNA  NM_001251 Homo sapiens CD68 antigen (CD68), mRNA  NM_000074 Homo sapiens tumor necrosis factor (ligand) superfamily, member 5 (hy syndrome) (TNFSF5), mRNA  NM_000073 Homo sapiens CD3G antigen, gamma polypeptide (TiT3 complex) (CD mRNA)	
NM_000747 Homo sapiens cholinergic receptor, nicotinic, beta polypeptide 1 (muscle (CHRNB1), mRNA  NM_000079 Homo sapiens cholinergic receptor, nicotinic, alpha polypeptide 1 (muscle (CHRNA1), mRNA  NM_001273 Homo sapiens chromodomain helicase DNA binding protein 4 (CHD4), NM_001271 Homo sapiens chromodomain helicase DNA binding protein 2 (CHD2), NM_001270 Homo sapiens chromodomain helicase DNA binding protein 1 (CHD1), NM_000078 Homo sapiens cholesteryl ester transfer protein, plasma (CETP), mRNA  NM_000076 Homo sapiens cyclin-dependent kinase inhibitor 1C (p57, Kip2) (CDKN mRNA  NM_001258 Homo sapiens cyclin-dependent kinase 3 (CDK3), mRNA  NM_001251 Homo sapiens CD68 antigen (CD68), mRNA  NM_000074 Homo sapiens tumor necrosis factor (ligand) superfamily, member 5 (hy syndrome) (TNFSF5), mRNA  NM_000073 Homo sapiens CD3G antigen, gamma polypeptide (TiT3 complex) (CD mRNA)	)),
(CHRNA1), mRNA  NM_001273 Homo sapiens chromodomain helicase DNA binding protein 4 (CHD4), NM_001271 Homo sapiens chromodomain helicase DNA binding protein 2 (CHD2), NM_001270 Homo sapiens chromodomain helicase DNA binding protein 1 (CHD1), NM_000078 Homo sapiens cholesteryl ester transfer protein, plasma (CETP), mRNA NM_000076 Homo sapiens cyclin-dependent kinase inhibitor 1C (p57, Kip2) (CDKN mRNA  NM_001258 Homo sapiens cyclin-dependent kinase 3 (CDK3), mRNA NM_001251 Homo sapiens CD68 antigen (CD68), mRNA  NM_000074 Homo sapiens tumor necrosis factor (ligand) superfamily, member 5 (hy syndrome) (TNFSF5), mRNA  NM_000073 Homo sapiens CD3G antigen, gamma polypeptide (TiT3 complex) (CD mRNA	<del>)</del>
NM 001271 Homo sapiens chromodomain helicase DNA binding protein 2 (CHD2), NM_001270 Homo sapiens chromodomain helicase DNA binding protein 1 (CHD1), NM_000078 Homo sapiens cholesteryl ester transfer protein, plasma (CETP), mRNA NM_000076 Homo sapiens cyclin-dependent kinase inhibitor 1C (p57, Kip2) (CDKN mRNA  NM_001258 Homo sapiens cyclin-dependent kinase 3 (CDK3), mRNA  NM_001251 Homo sapiens CD68 antigen (CD68), mRNA  NM_000074 Homo sapiens tumor necrosis factor (ligand) superfamily, member 5 (hy syndrome) (TNFSF5), mRNA  NM_000073 Homo sapiens CD3G antigen, gamma polypeptide (TiT3 complex) (CD mRNA	
NM_001270Homo sapiens chromodomain helicase DNA binding protein 1 (CHD1), NM_000078Homo sapiens cholesteryl ester transfer protein, plasma (CETP), mRNANM_000076Homo sapiens cyclin-dependent kinase inhibitor 1C (p57, Kip2) (CDKN mRNANM_001258Homo sapiens cyclin-dependent kinase 3 (CDK3), mRNANM_001251Homo sapiens CD68 antigen (CD68), mRNANM_000074Homo sapiens tumor necrosis factor (ligand) superfamily, member 5 (hy syndrome) (TNFSF5), mRNANM_000073Homo sapiens CD3G antigen, gamma polypeptide (TiT3 complex) (CD mRNA	mRNA
NM_001270Homo sapiens chromodomain helicase DNA binding protein 1 (CHD1), NM_000078Homo sapiens cholesteryl ester transfer protein, plasma (CETP), mRNANM_000076Homo sapiens cyclin-dependent kinase inhibitor 1C (p57, Kip2) (CDKN mRNANM_001258Homo sapiens cyclin-dependent kinase 3 (CDK3), mRNANM_001251Homo sapiens CD68 antigen (CD68), mRNANM_000074Homo sapiens tumor necrosis factor (ligand) superfamily, member 5 (hy syndrome) (TNFSF5), mRNANM_000073Homo sapiens CD3G antigen, gamma polypeptide (TiT3 complex) (CD mRNA	mRNA
NM_000078Homo sapiens cholesteryl ester transfer protein, plasma (CETP), mRNANM_000076Homo sapiens cyclin-dependent kinase inhibitor 1C (p57, Kip2) (CDKN mRNA)NM_001258Homo sapiens cyclin-dependent kinase 3 (CDK3), mRNANM_001251Homo sapiens CD68 antigen (CD68), mRNANM_000074Homo sapiens tumor necrosis factor (ligand) superfamily, member 5 (hy syndrome) (TNFSF5), mRNANM_000073Homo sapiens CD3G antigen, gamma polypeptide (TiT3 complex) (CD mRNA)	mRNA .
mRNA  NM_001258 Homo sapiens cyclin-dependent kinase 3 (CDK3), mRNA  NM_001251 Homo sapiens CD68 antigen (CD68), mRNA  NM_000074 Homo sapiens tumor necrosis factor (ligand) superfamily, member 5 (hy syndrome) (TNFSF5), mRNA  NM_000073 Homo sapiens CD3G antigen, gamma polypeptide (TiT3 complex) (CD mRNA	
NM_001251Homo sapiens CD68 antigen (CD68), mRNANM_000074Homo sapiens tumor necrosis factor (ligand) superfamily, member 5 (hy syndrome) (TNFSF5), mRNANM_000073Homo sapiens CD3G antigen, gamma polypeptide (TiT3 complex) (CD mRNA	(1C),
NM_001251Homo sapiens CD68 antigen (CD68), mRNANM_000074Homo sapiens tumor necrosis factor (ligand) superfamily, member 5 (hy syndrome) (TNFSF5), mRNANM_000073Homo sapiens CD3G antigen, gamma polypeptide (TiT3 complex) (CD mRNA	
NM_000074 Homo sapiens tumor necrosis factor (ligand) superfamily, member 5 (hy syndrome) (TNFSF5), mRNA  NM_000073 Homo sapiens CD3G antigen, gamma polypeptide (TiT3 complex) (CD mRNA	
mRNA	
	3G),
mRNA	PD5),
NM_001248 Homo sapiens ectonucleoside triphosphate diphosphohydrolase 3 (ENT: mRNA	PD3),
NM_001246 Homo sapiens ectonucleoside triphosphate diphosphohydrolase 2 (ENT mRNA	PD2),
NM_000072 Homo sapiens CD36 antigen (collagen type I receptor, thrombospondin (CD36), mRNA	receptor)
NM 000591 Homo sapiens CD14 antigen (CD14), mRNA	
NM 000071 Homo sapiens cystathionine-beta-synthase (CBS), mRNA	
NM 000388 Homo sapiens calcium-sensing receptor (hypocalciuric hypercalcemia 1	
neonatal hyperparathyroidism) (CASR), mRNA	, severe
NM 000070 Homo sapiens calpain 3, (p94) (CAPN3), mRNA	, severe
NM_000069 Homo sapiens calcium channel, voltage-dependent, L type, alpha 1S su (CACNA1S), mRNA	, severe
NM 001215 Homo sapiens carbonic anhydrase VI (CA6), mRNA	
NM 000067 Homo sapiens carbonic anhydrase II (CA2), mRNA	

	a la martida (CSG) mPNA
NM_000606	Homo sapiens complement component 8, gamma polypeptide (C8G), mRNA
NM 000066	Homo sapiens complement component 8, beta polypeptide (C8B), mRNA  Homo sapiens complement component 8, beta polypeptide (C8A), mRNA
NM_000562	Homo sapiens complement component 8, alpha polypeptide (C8A), mRNA  Homo sapiens complement component 8, alpha polypeptide (C8A), mRNA
VM 000587	Homo sapiens complement component 7 (C7), mRNA
NM 000064	Homo sapiens complement component 3 (C3), mRNA  Homo sapiens complement component 3 (C3), mRNA
	The Destan agammaglobulinemia TVTOSINE KIIIase (DIK), IIId II
NIM 001206	Homo sapiens basic transcription element binding protein 1 (B1EB1), index1
= = 000000	II remiens histinidase (RTD) mRNA
NM 001201	Homo seniens hone morphogenetic protein 3 (osteogenic) (BIMF3), IMNIVA
NM 001200	Homo saniens hone morphogenetic protein 2 (BMP2), mkNA
NM 000386	Homo sapiens bleomycin hydrolase (BLMH), mRNA
NM 000057	II serious Placem syndrome (BLM), mKNA
NM 001198	Home conjune PR domain containing I, with ZNF dolland (1 KDWI), mid-
NM 001196	TI Described DU2 interacting domain death agonist (BID), IIINNA
NM_000056	the household chain keto acid dehydrogenase E1, beta polypeptide
14141_000050	(maple syrup urine disease) (BCKDHB), nuclear gene encoding mitochondrial
	The state of the s
NM 000465	TI DESCRIPTION OF THE PROPERTY
NM 000705	TT ATPage H+/K+ exchanging, beta polypeptide (ATT4D), indivi-
NM_000049	Homo sapiens aspartoacylase (aminoacylase 2, Canavan disease) (ASPA),
14141_000045	mRNA
NM_000046	116 to an D (ADSR) mRNA
NM_000639	Homo sapiens aryisulfatase B (ARSB), interfer Homo sapiens tumor necrosis factor (ligand) superfamily, member 6 (TNFSF6),
14141_000055	DATA
NM_000042	Homo sapiens apolipoprotein H (beta-2-glycoprotein I) (APOH), mRNA
NM 000042	Homo sapiens apolipoprotein E (APOE), mRNA
NM 000040	Homo saniens anolipoprotein C-III (APOC3), mRNA
NM 000039	Homo saniens anolipoprotein A-I (APOAI), mRNA
NM 000038	Homo saniens adenomatosis polyposis coli (APC), mknA
NM 001157	Homo sapiens annexin A11 (ANXA11), mRNA
NM 001147	Try provides angionojetin 2 (ANGPTZ), mRNA
NM 001147	Transfer of the state of the st
NM_000036	Homo sapiens angiogenini, ribonaciease, retrassitation (AMPD1), Homo sapiens adenosine monophosphate deaminase 1 (isoform M) (AMPD1),
MM_000030	DATA
NM 001141	Home saniers arachidonate 15-lipoxygenase, second type (ALOXISB), mknA
	Home conjens aldolase B fructose-bisphosphate (ALDOB), higher
NM_000035	III remiens oldologe A fructose-hisphosphate (ALDOA), hikina
NM 000034	TT one ominolexalinate delta- synthase 2 (sidelobiasiie/ilypoetholino
NM_000032	· ) (AT ACC) mustor gene encoding mitochondral protein, include
273 4 000020	Trans agricus alanine alvoyulate aminotransierase (Oxalosis i, hyperoxalula i,
NM_000030	glycolicaciduria: serine-pyruvate aminotransierase) (AOA1), midva
NM 001126	Homo seniens adenylosuccinate synthase (ADSS), IIINVA
	Home caniens adrenergic heta-1-, receptor (ADRB1), likiva
NM_000684	Homo sapiens ADP-ribosylarginine hydrolase (ADPRH), mRNA
NM_001125	Homo sapiens adenylate cyclase 9 (ADCY9), mRNA
NM_001116	Homo sapiens adenylate cyclase 8 (brain) (ADCY8), mRNA
NM 001115	Try and adamylets evoluse 7 (ADCY/) MKNA
NM_001114	The recognitions a disjute or and metalloproteinase domain & (ADAMO), Illicity
NM_001109	Homo sapiens a disintegrin and metalloproteinase domain 10 (ADAM10),
NM_001110	HOMO Sapiens a disintegrin and members and
37.6 001100	mRNA Homo sapiens acylphosphatase 2, muscle type (ACYP2), mRNA
NM_001108	
NM_001107	Homo sapiens acyiphosphaiase i, cryunocyte (com-

	The state of the s
NM_001104	Homo sapiens actinin, alpha 3 (ACTN3), mRNA
NM_001086	Homo sapiens arylacetamide deacetylase (esterase) (AADAC), mRNA
NM_001043	Homo sapiens solute carrier family 6 (neurotransmitter transporter,
	noradrenalin), member 2 (SLC6A2), mRNA
NM_000532	Homo sapiens propionyl Coenzyme A carboxylase, beta polypeptide (PCCB),
17.5.000770	nuclear gene encoding mitochondrial protein, mRNA
NM_002579	Homo sapiens paralemmin (PALM), mRNA
NM_002443	Homo sapiens microseminoprotein, beta- (MSMB), mRNA
NM_002418	Homo sapiens motilin (MLN), mRNA
NM_002300	Homo sapiens lactate dehydrogenase B (LDHB), mRNA
NM_002243	Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 15
NR 6 001504	(KCNJ15), mRNA
NM_001534	Homo sapiens homeo box 11-like 1 (HOX11L1), mRNA
NM_001454	Homo sapiens forkhead box J1 (FOXJ1), mRNA
NM_004001	Homo sapiens Fc fragment of IgG, low affinity IIb, receptor for (CD32) (FCGR2B), mRNA
NM_001276	Homo sapiens chitinase 3-like 1 (cartilage glycoprotein-39) (CHI3L1), mRNA
NM_001752	Homo sapiens catalase (CAT), mRNA
NM_001610	Homo sapiens acid phosphatase 2, lysosomal (ACP2), mRNA
NM_003461	Homo sapiens zyxin (ZYX), mRNA
NM_003460	Homo sapiens zona pellucida glycoprotein 2 (sperm receptor) (ZP2), mRNA
NM_003459	Homo sapiens solute carrier family 30 (zinc transporter), member 3 (SLC30A3),
	mRNA
NM_003430	Homo sapiens zinc finger protein 91 (HPF7, HTF10) (ZNF91), mRNA
NM_003429	Homo sapiens zinc finger protein 85 (HPF4, HTF1) (ZNF85), mRNA
NM_003428	Homo sapiens zinc finger protein 84 (HPF2) (ZNF84), mRNA
NM_003416	Homo sapiens zinc finger protein 7 (KOX 4, clone HF.16) (ZNF7), mRNA
NM_003427	Homo sapiens zinc finger protein 76 (expressed in testis) (ZNF76), mRNA
NM_003426	Homo sapiens zinc finger protein 74 (Cos52) (ZNF74), mRNA
NM_003425	Homo sapiens zinc finger protein 45 (a Kruppel-associated box (KRAB) domain
	polypeptide) (ZNF45), mRNA
NM_003423	Homo sapiens zinc finger protein 43 (HTF6) (ZNF43), mRNA
NM_003422	Homo sapiens zinc finger protein 42 (myeloid-specific retinoic acid- responsive) (ZNF42), mRNA
NM 003420	Homo sapiens zinc finger protein 35 (clone HF.10) (ZNF35), mRNA
NM 003458	Homò sapiens bassoon (presynaptic cytomatrix protein) (BSN), mRNA
NM_003456	Homo sapiens zinc finger protein 205 (ZNF205), mRNA
NM 003453	Homo sapiens zinc finger protein 198 (ZNF198), mRNA
NM 003450	Homo sapiens zinc finger protein 174 (ZNF174), mRNA
NM 003447	Homo sapiens zinc finger protein 165 (ZNF165), mRNA
NM 003446	Homo sapiens zinc finger protein 157 (HZF22) (ZNF157), mRNA
NM 003443	Homo sapiens zinc finger protein 151 (pHZ-67) (ZNF151), mRNA
NM 003442	Homo sapiens zinc finger protein 143 (clone pHZ-1) (ZNF143), mRNA
NM 003441	Homo sapiens zinc finger protein 141 (clone pHZ-44) (ZNF141), mRNA
NM 003440	Homo sapiens zinc finger protein 140 (clone pHZ-39) (ZNF140), mRNA
NM 003438	Homo sapiens zinc finger protein 137 (clone pHZ-30) (ZNF137), mRNA
NM 003437	Homo sapiens zinc finger protein 136 (clone pHZ-20) (ZNF136), mRNA
NM 003436	Homo sapiens zinc finger protein 135 (clone pHZ-17) (ZNF135), mRNA
NM 003435	Homo sapiens zinc finger protein 134 (clone pHZ-15) (ZNF134), mRNA
NM 003434	Homo sapiens zinc finger protein 133 (clone pHZ-13) (ZNF133), mRNA
NM 003433	Homo sapiens zinc finger protein 132 (clone pHZ-12) (ZNF132), mRNA
NM 003431	Homo sapiens zinc finger protein 124 (HZF-16) (ZNF124), mRNA
11112 005 151	1

	TILLA (TEV) mPNA
NM 003411	Homo sapiens zinc finger protein, Y-linked (ZFY), mRNA
373 £ 002 410	Home conjent zinc finger protein X-linked (ZFA), IIIKNA
NM 003405	Homo sapiens tyrosine 3-monooxygenase/tryptophan 5-monooxygenase
	estivation protein, eta polypentide (YWHAH), mKNA
NIM 003404	Homo sapiens tyrosine 3-monooxygenase/tryptopnan 3-monooxygenase
	- timetical protein, hete polymentide (YWHAB), MKNA
NM 000380	Homo sapiens xeroderma pigmentosum, complementation group A (XPA),
14141_000000	mPNA
NM 003931	Homo saniens WAS protein family, member 1 (WASF1), mRNA
NM 003384	Theme conjugate vaccinia related kinase I (VRKI), mKNA
NM 003383	Home conjone very low density linoprotein receptor (VLDLR), IIIRNA
NM 003383	Homo sapiens vasoactive intestinal peptide receptor 2 (VIPR2), mRNA
	Homo sapiens vasoactive intestinal peptide (VIP), mRNA
NM_003381	Homo sapiens vimentin (VIM), mRNA
NM_003380	Homo sapiens vascular endothelial growth factor B (VEGFB), mRNA
NM_003377	Homo sapiens vascular endothelial growth factor (VEGF), mRNA  Homo sapiens vascular endothelial growth factor (VEGF), mRNA
NM_003376	Homo sapiens viscular endomenal growth factor (D3) receptor (VDR), mRNA  Homo sapiens vitamin D (1,25- dihydroxyvitamin D3) receptor (VDR), mRNA
NM_000376	Homo sapiens vitamin D (1,25- diliydroxy vitamini B5) receptor (1,25), mRNA
NM_003375	Homo sapiens voltage-dependent anion channel 2 (VDAC2), mRNA  Homo sapiens voltage-dependent anion channel 2 (VDAC1), mRNA
NM_003374	Homo sapiens voltage-dependent anion channel 1 (VDAC1), mRNA
NM 003371	Homo sapiens vav 2 oncogene (VAV2), mRNA
NM 003370	Homo sapiens vasodilator-stimulated phosphoprotein (VASP), mRNA
NM 003762	Homo sapiens vesicle-associated membrane protein 4 (VAMP4), mRNA  (MMPAC) - PRNA
NM 003369	There agricus ITV radiation resistance associated gene (UVRAU), IIIRNA
NM 003577	Homo sapiens undifferentiated embryonic cell transcription factor 1 (UTF1),
	1 TONIA
NM_003470	Homo sapiens ubiquitin specific protease 7 (herpes virus-associated) (USP7),
	DNIA
NM 003481	Homo caniens ubiquitin specific protease 5 (isopeptidase T) (USP3), mRNA
NM 003363	Homo saniens ubiquitin specific protease 4 (proto-oncogene) (USF4), micha
NM 003368	Trans geniene ubiquitin specific protease 1 (USP1), mRNA
NM 003940	TI-me senions ubiquitin specific protease 13 (ISODEDIIGASE 1-3) (USF 13), IIICNA
NM_003367	I Home ganiens unstream transcription factor 2, c-tos interacting (USF2), IIICNA
NM_003366	Homo sapiens ubiquinol-cytochrome c reductase core protein II (UQCRC2),
MM_002200	mp D N A
ND 6 002265	Homo sapiens ubiquinol-cytochrome c reductase core protein I (UQCRC1),
NM_003365	
	mRNA  Homo sapiens uridine phosphorylase (UP), mRNA  [MOD]
NM_003364	Homo sapiens uridine phosphorylase (OI), med 12  Homo sapiens uromodulin (uromucoid, Tamm-Horsfall glycoprotein) (UMOD),
NM_003361	
	mRNA Homo sapiens Kruppel-like factor 7 (ubiquitous) (KLF7), mRNA
NM_003709	Homo sapiens Kruppei-like lactor / (dolydrous) (12277); http://www.forese.g. Q. IDP-galactose ceramide
NM_003360	Homo sapiens UDP glycosyltransferase 8 (UDP-galactose ceramide
	galactosyltransferase) (UGT8), mRNA
NM_001074	Homo sapiens UDP glycosyltransferase 2 family, polypeptide B7 (UGT2B7),
	mRNA Comiliar polymontide R17 (IGT2R17)
NM_001077	Homo sapiens UDP glycosyltransferase 2 family, polypeptide B17 (UGT2B17),
	mRNA 0. S. wiles polymontide R15 (LIGT2R15)
NM_001076	Homo sapiens UDP glycosyltransferase 2 family, polypeptide B15 (UGT2B15),
_	DNTA
	T : rmp 1
NM 001075	Homo sapiens UDP glycosyltransierase 2 family, polypeptide 216 (C G 222 23)
NM_001075	mRNA
NM_001075 NM_003359	Homo sapiens UDP glycosyltransferase 2 family, polypeptide B10 (UGT2B10), mRNA  Homo sapiens UDP-glucose dehydrogenase (UGDH), mRNA  Homo sapiens UDP-glucose ceramide glucosyltransferase (UGCG), mRNA

	11' GIOD) DNA
NM_003357	Homo sapiens uteroglobin (UGB), mRNA
NM_003352	Homo sapiens ubiquitin-like 1 (sentrin) (UBL1), mRNA
NM_003347	Homo sapiens ubiquitin-conjugating enzyme E2L 3 (UBE2L3), mRNA
NM_003337	Homo sapiens ubiquitin-conjugating enzyme E2B (RAD6 homolog) (UBE2B),
	mRNA
NM_003336	Homo sapiens ubiquitin-conjugating enzyme E2A (RAD6 homolog) (UBE2A),
	mRNA
NM_003335	Homo sapiens ubiquitin-activating enzyme E1-like (UBE1L), mRNA
NM_000550	Homo sapiens tyrosinase-related protein 1 (TYRP1), mRNA
NM_000372	Homo sapiens tyrosinase (oculocutaneous albinism IA) (TYR), mRNA
NM_001071	Homo sapiens thymidylate synthetase (TYMS), mRNA
NM_003331	Homo sapiens tyrosine kinase 2 (TYK2), mRNA
NM_003330	Homo sapiens thioredoxin reductase 1 (TXNRD1), mRNA
NM_003329	Homo sapiens thioredoxin (TXN), mRNA
NM_003328	Homo sapiens TXK tyrosine kinase (TXK), mRNA
NM_003324	Homo sapiens tubby like protein 3 (TULP3), mRNA
NM 003323	Homo sapiens tubby like protein 2 (TULP2), mRNA
NM_003321	Homo sapiens Tu translation elongation factor, mitochondrial (TUFM), mRNA
NM_001070	Homo sapiens tubulin, gamma 1 (TUBG1), mRNA
NM_001069	Homo sapiens tubulin, beta polypeptide (TUBB), mRNA
NM_000371	Homo sapiens transthyretin (prealbumin, amyloidosis type I) (TTR), mRNA
NM 000370	Homo sapiens tocopherol (alpha) transfer protein (ataxia (Friedreich-like) with
	vitamin E deficiency) (TTPA), mRNA
NM_003319	Homo sapiens titin (TTN), mRNA
NM 003318	Homo sapiens TTK protein kinase (TTK), mRNA
NM 003317	Homo sapiens thyroid transcription factor 1 (TITF1), mRNA
NM_003315	Homo sapiens tetratricopeptide repeat domain 2 (TTC2), mRNA
NM_003314	Homo sapiens tetratricopeptide repeat domain 1 (TTC1), mRNA
NM_003311	Homo sapiens tumor suppressing subtransferable candidate 3 (TSSC3), mRNA
NM_003310	Homo sapiens tumor suppressing subtransferable candidate 1 (TSSC1), mRNA
NM_000369	Homo sapiens thyroid stimulating hormone receptor (TSHR), mRNA
NM_000549	Homo sapiens thyroid stimulating hormone, beta (TSHB), mRNA
NM_003496	Homo sapiens transformation/transcription domain-associated protein (TRRAP),
NM 003301	mRNA Homo sapiens thyrotropin-releasing hormone receptor (TRHR), mRNA
NM 003299	Homo sapiens tumor rejection antigen (gp96) 1 (TRA1), mRNA
	Homo sapiens nuclear receptor subfamily 2, group C, member 2 (NR2C2),
NM_003298	mRNA
NM_003296	Homo sapiens testis specific protein 1 (probe H4-1 p3-1) (TPX1), mRNA
NM_003295	Homo sapiens tumor protein, translationally-controlled 1 (TPT1), mRNA
NM_003595	Homo sapiens tyrosylprotein sulfotransferase 2 (TPST2), mRNA
NM_003292	Homo sapiens translocated promoter region (to activated MET oncogene) (TPR),
) D. C. 0000001	mRNA
NM_003291	Homo sapiens tripeptidyl peptidase II (TPP2), mRNA
NM_000547	Homo sapiens thyroid peroxidase (TPO), nuclear gene encoding mitochondrial protein, mRNA
NM 003290	Homo sapiens tropomyosin 4 (TPM4), mRNA
NM_003289	Homo sapiens tropomyosin 2 (beta) (TPM2), mRNA
NM_000366	Homo sapiens tropomyosin 1 (alpha) (TPM1), mRNA
NM_000365	Homo sapiens triosephosphate isomerase 1 (TPI1), mRNA
NM 003288	Homo sapiens tumor protein D52-like 2 (TPD52L2), mRNA
NM_003288	Homo sapiens tumor protein D52-like 1 (TPD52L1), mRNA
1414 003207	1 Home suprem tumor protein DSE-like I (11 DSED1), Hillers

	TODAY MILLA (TODAD) mPNA
NM_003935	Homo sapiens topoisomerase (DNA) III beta (TOP3B), mRNA
NM_001067	Homo sapiens topoisomerase (DNA) II alpha (170kD) (TOP2A), mRNA
NM_003285	Homo sapiens tenascin R (restrictin, janusin) (TNR), mRNA
NM_003284	Homo sapiens transition protein 1 (during histone to protamine replacement)
_	(TNP1), mRNA
NM 000364	Homo sapiens troponin T2, cardiac (TNNT2), mRNA
NM_003283	Homo sapiens troponin T1, skeletal, slow (TNNT1), mRNA
NM 000363	Homo saniens trononin I. cardiac (TNNI3), mRNA
NM 003282	Homo sapiens troponin I, skeletal, fast (TNNI2), mRNA
NM 003281	Homo sapiens troponin I, skeletal, slow (TNNII), mRNA
NM 003279	Homo saniens troponin C2, fast (TNNC2), mRNA
NM 003280	Homo saniens troponin C. slow (TNNC1), mRNA
NM 003280	Transport tracing kingse non-recentor, I (INKI), IIIKNA
	Homo sapiens tyrosine kinase, non receptor, 1 (1997) Homo sapiens tumor necrosis factor (ligand) superfamily, member 8 (TNFSF8),
NM_001244	DATA
377 £ 001050	Homo sapiens tumor necrosis factor (ligand) superfamily, member 7 (TNFSF7),
NM_001252	
-7.6.000006	mRNA Homo sapiens tumor necrosis factor (ligand) superfamily, member 4 (tax-
NM_003326	transcriptionally activated glycoprotein 1, 34kD) (TNFSF4), mRNA
	Homo sapiens tumor necrosis factor (ligand) superfamily, member 13
NM_003808	Homo sapiens tuinor necrosis factor (figure) superior
	(TNFSF13), mRNA  Homo sapiens tumor necrosis factor (ligand) superfamily, member 12
NM_003809	Homo sapiens tumor necrosis factor (figure) superformity, internet
	(TNFSF12), mRNA  Homo sapiens tumor necrosis factor (ligand) superfamily, member 10
NM_003810	Homo sapiens tumor necrosis factor (figalia) superfamily, member 19
L	(TNFSF10), mRNA
NM_001243	Homo sapiens tumor necrosis factor receptor superfamily, member 8
	(TNFRSF8), mRNA
NM_001242	Homo sapiens tumor necrosis factor receptor superfamily, member 7
	(TNFRSF7), mRNA
NM_000043	Homo sapiens tumor necrosis factor receptor superfamily, member 6
	(TNFRSF6), mRNA
NM 003327	Homo sapiens tumor necrosis factor receptor superfamily, member 4
_	(TNIED SEA) mDNA
NM 001066	Homo sapiens tumor necrosis factor receptor superfamily, member 1B
_	(TNEDSEIR) mRNA
NM 001065	Homo sapiens tumor necrosis factor receptor superfamily, member 1A
1,11,2_0	(TNEP SEI A) mRNA
NM_001192	Homo sapiens tumor necrosis factor receptor superfamily, member 17
1414_001152	(TNIED SE17) mRNA
NM 003820	Homo sapiens tumor necrosis factor receptor superfamily, member 14
14141_003020	Chernesvirus entry mediator) (TNFRSF14), mKINA
NIM 003790	Home seniers tymer necrosis factor receptor superfamily, member 12
NM_003790	(translocating chain-association membrane protein) (INFRSF12), mkNA
NTM 002546	Homo sapiens tumor necrosis factor receptor superfamily, member 11b
NM_002546	(coteoprotegerin) (TNFRSF11R) mRNA
ND 6 0000000	Homo sapiens tumor necrosis factor receptor superfamily, member 11a, activate
NM_003839	AFNIEKB (TNIERSEILA) mRNA
37 C 0000 40	Homo sapiens tumor necrosis factor receptor superfamily, member 10d, decoy
NM_003840	with truncated death domain (TNFRSF10D), mRNA
NM_003842	HOMO Sapiens lumor necrosis factor receptor superfamily, memor 135
	(TNFRSF10B), mRNA  Homo sapiens tumor necrosis factor receptor superfamily, member 10a
NM 003844	Homo sapiens tumor necrosis factor receptor superfamily, member for

	(TRIED OF 10 A) DAIA
77.5.000076	(TNFRSF10A), mRNA
NM_003276	Homo sapiens thymopoietin (TMPO), mRNA
NM_003275	Homo sapiens tropomodulin (TMOD), mRNA
NM_003274	Homo sapiens transmembrane protein 1 (TMEM1), mRNA
NM_003692	Homo sapiens transmembrane protein with EGF-like and two follistatin-like
27.6.000072	domains 1 (TMEFF1), mRNA  Homo sapiens transmembrane 7 superfamily member 2 (TM7SF2), mRNA
NM_003273	Homo sapiens transmemorane 7 superfamily member 2 (1147312), interval  Homo sapiens transmembrane 7 superfamily member 1 (upregulated in kidney)
NM_003272	(TM7SF1), mRNA
NM_003271	Homo sapiens transmembrane 4 superfamily member 7 (TM4SF7), mRNA
NM 003270	Homo sapiens transmembrane 4 superfamily member 6 (TM4SF6), mRNA
NM 003963	Homo sapiens transmembrane 4 superfamily member 5 (TM4SF5), mRNA
NM_003269	Homo sapiens nuclear receptor subfamily 2, group E, member 1 (NR2E1), mRNA
NM 003266	Homo sapiens toll-like receptor 4 (TLR4), mRNA
NM 003265	Homo sapiens toll-like receptor 3 (TLR3), mRNA
NM 003264	Homo sapiens toll-like receptor 2 (TLR2), mRNA
NM 003263	Homo sapiens toll-like receptor 1 (TLR1), mRNA
NM 003258	Homo sapiens thymidine kinase 1, soluble (TK1), mRNA
NM 003257	Homo sapiens tight junction protein 1 (zona occludens 1) (TJP1), mRNA
NM 003256	Homo sapiens tissue inhibitor of metalloproteinase 4 (TIMP4), mRNA
NM 003254	Homo sapiens tissue inhibitor of metalloproteinase 1 (erythroid potentiating
11111_005251	activity, collagenase inhibitor) (TIMP1), mRNA
NM 003597	Homo sapiens TGFB inducible early growth response 2 (TIEG2), mRNA
NM 003253	Homo sapiens T-cell lymphoma invasion and metastasis 1 (TIAM1), mRNA
NM 000460	Homo sapiens thrombopoietin (myeloproliferative leukemia virus oncogene
	ligand, megakaryocyte growth and development factor) (THPO), mRNA
NM 003249	Homo sapiens thimet oligopeptidase 1 (THOP1), mRNA
NM 003248	Homo sapiens thrombospondin 4 (THBS4), mRNA
NM 003247	Homo sapiens thrombospondin 2 (THBS2), mRNA
NM 003246	Homo sapiens thrombospondin 1 (THBS1), mRNA
NM 000361	Homo sapiens thrombomodulin (THBD), mRNA
NM 000360	Homo sapiens tyrosine hydroxylase (TH), mRNA
NM 003241	Homo sapiens transglutaminase 4 (prostate) (TGM4), mRNA
NM_003245	Homo sapiens transglutaminase 3 (E polypeptide, protein-glutamine-gamma-glutamyltransferase) (TGM3), mRNA
NM 000359	Homo sapiens transglutaminase 1 (K polypeptide epidermal type I, protein-
14147_000223	glutamine-gamma-glutamyltransferase) (TGM1), mRNA
NM 003243	Homo sapiens transforming growth factor, beta receptor III (betaglycan, 300kD)
14141_003243	(TGFBR3), mRNA
NM 003242	Homo sapiens transforming growth factor, beta receptor II (70-80kD)
1111_003272	(TGFBR2), mRNA
NM 000358	Homo sapiens transforming growth factor, beta-induced, 68kD (TGFBI), mRNA
NM 003239	Homo sapiens transforming growth factor, beta 3 (TGFB3), mRNA
NM 003238	Homo sapiens transforming growth factor, beta 2 (TGFB2), mRNA
NM 003236	Homo sapiens transforming growth factor, alpha (TGFA), mRNA
NM 003234	Homo sapiens transferrin receptor (p90, CD71) (TFRC), mRNA
NM 003237	Homo sapiens transferrin receptor 2 (TFR2), mRNA
NM 003227	Homo sapiens trefoil factor 3 (intestinal) (TFF3), mRNA
NM 003225	Homo sapiens trefoil factor 1 (breast cancer, estrogen-inducible sequence
	expressed in) (TFF1), mRNA
NM_003224	Homo sapiens ADP-ribosylation factor related protein 1 (ARFRP1), mRNA

	· · · · · · · · · · · · · · · · · · ·
NM_003219	Homo sapiens telomerase reverse transcriptase (TERT), mRNA
NM_003673	Homo sapiens titin-cap (telethonin) (TCAP), mRNA
NM_003217	Homo sapiens testis enhanced gene transcript (TEGT), mRNA
NM_003216	Homo sapiens thyrotrophic embryonic factor (TEF), mRNA
NM_003213	Homo sapiens TEA domain family member 4 (TEAD4), mRNA
NM_003211_	Homo sapiens thymine-DNA glycosylase (TDG), mRNA
NM_003608	Homo sapiens G protein-coupled receptor 65 (GPR65), mRNA
NM_000355	Homo sapiens transcobalamin II; macrocytic anemia (TCN2), mRNA
NM_001062	Homo sapiens transcobalamin I (vitamin B12 binding protein, R binder family) (TCN1), mRNA
NM 003202	Homo saniens transcription factor 7 (T-cell specific, HMG-box) (TCF7), mRNA
NM_003201	Homo sapiens transcription factor 6-like 1 (mitochondrial transcription factor 1-like) (TCF6L1), mRNA
NM 003199	Homo sapiens transcription factor 4 (TCF4), mRNA
NM_003206	Homo sapiens transcription factor 21 (TCF21), mRNA
NM_000545	Homo sapiens transcription factor 1, hepatic; LF-B1, hepatic nuclear factor
NR 6 000100	(HNF1), albumin proximal factor (TCF1), mRNA  Homo sapiens transcription elongation factor B (SIII), polypeptide 3 (110kD,
NM_003198	Homo sapiens transcription clongation factor is (Sin), polypopulae's (Toke),
>D 6 001060	elongin A) (TCEB3), mRNA  Homo sapiens thromboxane A2 receptor (TBXA2R), mRNA
NM_001060	Homo sapiens TATA box binding protein (TBA), mRNA
NM_003194	Homo sapiens tubulin-specific chaperone c (TBCC), mRNA
NM_003192	Homo sapiens tafazzin (cardiomyopathy, dilated 3A (X-linked); endocardial
NM_000116	fibroelastosis 2; Barth syndrome) (TAZ), mRNA
NM_000353	Homo sapiens tyrosine aminotransferase (TAT), nuclear gene encoding
	mitochondrial protein, mRNA
NM 003191	Homo sapiens threonyl-tRNA synthetase (TARS), mRNA
NM 003190	Homo sapiens TAP binding protein (tapasin) (TAPBP), mRNA
NM 003189	Homo sapiens T-cell acute lymphocytic leukemia 1 (TAL1), mRNA
NM_003188	Homo sapiens mitogen-activated protein kinase kinase kinase 7 (MAP3K7), mRNA
NM_003487	Homo sapiens TATA box binding protein (TBP)-associated factor, RNA
1111_005 107	polymerase II, N, 68kD (RNA-binding protein 56) (TAF2N), mRNA
NM_003187	Homo sapiens TATA box binding protein (TBP)-associated factor, RNA
11112_00020.	polymerase II, G, 32kD (TAF2G), mRNA
NM 001057	Homo sapiens tachykinin receptor 2 (TACR2), mRNA
NM_003180	Homo sapiens synaptotagmin 5 (SYT5), mRNA
NM 003895	Homo sapiens synaptojanin 1 (SYNJ1), mRNA
NM 003490	Homo sapiens synapsin III (SYN3), mRNA
NM 003178	Homo sapiens synapsin II (SYN2), mRNA
NM 003177	Homo sapiens spleen tyrosine kinase (SYK), mRNA
NM 003176	Homo sapiens synaptonemal complex protein 1 (SYCP1), mRNA
NM 003172	Homo sapiens surfeit 1 (SURF1), mRNA
NM 003167	Homo sapiens sulfotransferase family, cytosolic, 2A, dehydroepiandrosterone
14117_005107	(DHEA) -preferring, member 1 (SULT2A1), mRNA
NM_001056	Homo sapiens sulfotransferase family, cytosolic, 1C, member 1 (SULT1C1),
1111_001000	mRNA
NM 001054	Homo sapiens sulfotransferase family, cytosolic, 1A, phenol-preferring, member
1111_001051	2 (SULT1A2), mRNA
NM_001055	Homo sapiens sulfotransferase family, cytosolic, 1A, phenol-preferring, member 1 (SULT1A1), mRNA
NM 003165	Homo sapiens syntaxin binding protein 1 (STXBP1), mRNA
TATAT 002 102	Tionio sapionis synakin omanig protein 1 (5112211), med 11

NM 003163	Homo sapiens syntaxin 1B (STX1B), mRNA
NM 003159	Homo sapiens serine/threonine kinase 9 (STK9), mRNA
NM 003158	Homo sapiens serine/threonine kinase 6 (STK6), mRNA
NM 003157	Homo sapiens serine/threonine kinase 2 (STK2), mRNA
NM 003600	Homo sapiens serine/threonine kinase 15 (STK15), mRNA
NM 003160	Homo sapiens serine/threonine kinase 13 (aurora/IPL1-like) (STK13), mRNA
NM 003156	Homo sapiens stromal interaction molecule 1 (STIM1), mRNA
NM 003155	Homo sapiens stanniocalcin 1 (STC1), mRNA
NM 003877	Homo sapiens STAT induced STAT inhibitor-2 (STATI2), mRNA
NM 003154	Homo sapiens statherin (STATH), mRNA
NM_003153	Homo sapiens signal transducer and activator of transcription 6, interleukin-4 induced (STAT6), mRNA
NM_003152	Homo sapiens signal transducer and activator of transcription 5A (STAT5A), mRNA
NM_003151	Homo sapiens signal transducer and activator of transcription 4 (STAT4), mRNA
NM_003150	Homo sapiens signal transducer and activator of transcription 3 (acute-phase response factor) (STAT3), mRNA
NM_000349	Homo sapiens steroidogenic acute regulatory protein (STAR), mRNA
NM_003473	Homo sapiens signal transducing adaptor molecule (SH3 domain and ITAM motif) 1 (STAM), mRNA
NM_003149	Homo sapiens src homology three (SH3) and cysteine rich domain (STAC), mRNA
NM_001048	Homo sapiens somatostatin (SST), mRNA
NM_003146	Homo sapiens structure specific recognition protein 1 (SSRP1), mRNA
NM_003745	Homo sapiens JAK binding protein (SSI-1), mRNA
NM_001080	Homo sapiens aldehyde dehydrogenase 5 family, member A1 (succinate-semialdehyde dehydrogenase) (ALDH5A1), mRNA
NM_003139	Homo sapiens signal recognition particle receptor ('docking protein') (SRPR), mRNA
NM_003138	Homo sapiens SFRS protein kinase 2 (SRPK2), mRNA
NM_003135	Homo sapiens signal recognition particle 19kD (SRP19), mRNA
NM_003132	Homo sapiens spermidine synthase (SRM), mRNA
NM_003130	Homo sapiens sorcin (SRI), mRNA
NM_001047	Homo sapiens steroid-5-alpha-reductase, alpha polypeptide 1 (3-oxo-5 alpha-steroid delta 4-dehydrogenase alpha 1) (SRD5A1), mRNA
NM_003743	Homo sapiens nuclear receptor coactivator 1 (NCOA1), mRNA
NM_003128	Homo sapiens spectrin, beta, non-erythrocytic 1 (SPTBN1), mRNA
NM_003127	Homo sapiens spectrin, alpha, non-erythrocytic 1 (alpha-fodrin) (SPTAN1), mRNA
NM_003126	Homo sapiens spectrin, alpha, erythrocytic 1 (elliptocytosis 2) (SPTA1), mRNA
NM_003125	Homo sapiens small proline-rich protein 1B (cornifin) (SPRR1B), mRNA
NM_003124	Homo sapiens sepiapterin reductase (7,8-dihydrobiopterin:NADP+ oxidoreductase) (SPR), mRNA
NM_003123	Homo sapiens sialophorin (gpL115, leukosialin, CD43) (SPN), mRNA
NM_003121	Homo sapiens Spi-B transcription factor (Spi-1/PU.1 related) (SPIB), mRNA
NM_003120	Homo sapiens spleen focus forming virus (SFFV) proviral integration oncogene spi1 (SPI1), mRNA
NM_003119	Homo sapiens spastic paraplegia 7, paraplegin (pure and complicated autosomal recessive) (SPG7), mRNA
NM_003118	Homo sapiens secreted protein, acidic, cysteine-rich (osteonectin) (SPARC), mRNA
NM_003112	Homo sapiens Sp4 transcription factor (SP4), mRNA

	was a Way (SOY4) mRNA
NM_003107	Homo sapiens SRY (sex determining region Y)-box 4 (SOX4), mRNA
NM_003108	Homo sapiens SRY (sex determining region Y)-box 11 (SOX11), mRNA
NM_003104	Homo sapiens sorbitol dehydrogenase (SORD), mRNA
NM_003102	Homo sapiens superoxide dismutase 3, extracellular (SOD3), mRNA
NM 003794	Homo sapiens sorting nexin 4 (SNX4), mRNA
NM 003100	Homo sapiens sorting nexin 2 (SNX2), mRNA
77.5.000004	There agains small nuclear ribonic leoprotein polypeptide E (SING E), midvi
NM 003092	Homo sapiens small nuclear ribonucleoprotein polypeptide B (SINKI B2),
	mRNA Homo sapiens small nuclear ribonucleoprotein polypeptide A' (SNRPA1),
NM_003090	DATA
NM_003089	Homo sapiens small nuclear ribonucleoprotein 70kD polypeptide (RNP antigen)
MM_002093	(SNRP70), mRNA
NTM 002409	II comions stennin (SNN) mRNA
NM_003498	Homo sapiens synuclein, gamma (breast cancer-specific protein 1) (SNCG),
NM_003087	Homo sapiens syndereni, guinna (oreast anni
	mRNA Homo sapiens small nuclear RNA activating complex, polypeptide 2, 45kD
NM_003083	Homo sapiens small nuclear KIVA activating company page 1
	(SNAPC2), mRNA Homo sapiens small nuclear RNA activating complex, polypeptide 1, 43kD
NM_003082	Homo sapiens small nuclear RNA activating complex, polypeptide 1, 1912
	(SNAPC1), mRNA
NM_003081	Homo sapiens synaptosomal-associated protein, 25kD (SNAP25), mRNA
NM_003078	Homo sapiens SWI/SNF related, matrix associated, actin dependent regulator of
_	chromatin, subfamily d, member 3 (SMARCD3), mRNA
NM_003077	Homo sapiens SWI/SNF related, matrix associated, actin dependent regulator of
_	1 1 mily d member 2 (SMARCD2), mKNA
NM_003076	Thomas aniens SWI/SNF related, matrix associated, actin dependent regulator of
11112_0000	1 1 months subfamily d. member 1 (SMARCD)), MKNA
NM_003075	Homo saniens SWI/SNF related, matrix associated, actiff dependent regulator of
14141_005075	1 1 marking subfamily a member 2 (SMARCU2), MKNA
NM_003074	Homo saniens SWI/SNF related, matrix associated, actin dependent regulator of
14141_003074	1 1 menting subfamily a member 1 (SMARCCI), MKNA
ND 4 002073	Homo sapiens SWI/SNF related, marria associated, actin dependent regulator of
NM_003073	1 . 1
77.5.000.601	Homo sapiens SWI/SNF related, matrix associated, actin dependent regulator of
NM_003601	chromatin, subfamily a, member 5 (SMARCA5), mRNA
	Homo sapiens SWI/SNF related, matrix associated, actin dependent regulator of
NM_003071	chromatin, subfamily a, member 3 (SMARCA3), mRNA
	Homo sapiens SWI/SNF related, matrix associated, actin dependent regulator of
NM_003070	Homo sapiens SWI/SNF related, matrix associated, actin dependent regulator
	chromatin, subfamily a, member 2 (SMARCA2), mRNA
NM_003069	Homo sapiens SWI/SNF related, matrix associated, actin dependent regulator of
	chromatin, subfamily a, member 1 (SMARCA1), mRNA
NM_003982	Homo sapiens solute carrier family 7 (cationic amino acid transporter, y+
	system), member 7 (SLC7A7), mRNA
NM 003046	Homo sapiens solute carrier family 7 (cationic amino acid transporter, y+
	system) member 2 (SLC7A2), mRNA
NM_003045	Homo sapiens solute carrier family 7 (cationic amino acid transporter, y+
14141_002042	austom) member 1 (SI C7A1) mRNA
NB 4 002042	Homo sapiens solute carrier family 6 (neurotransmitter transporter, taurine),
NM_003043	mambar 6 (SI C6A6) mRNA
NM_003043	
	Hemos serious solute carrier family 6 (neurotransmitter transporter, serotonin),
NM_003043	Homo sapiens solute carrier family 6 (neurotransmitter transporter, serotonin), member 4 (SLC6A4), mRNA

<del>.,</del>	o (or octo) DVI
	member 3 (SLC6A3), mRNA
NM_003042	Homo sapiens solute carrier family 6 (neurotransmitter transporter, GABA),
	member 1 (SLC6A1), mRNA
NM_003044	Homo sapiens solute carrier family 6 (neurotransmitter transporter,
	betaine/GABA), member 12 (SLC6A12), mRNA
NM_000453	Homo sapiens solute carrier family 5 (sodium iodide symporter), member 5
_	(SLC5A5), mRNA
NM_003041	Homo sapiens solute carrier family 5 (sodium/glucose cotransporter), member 2
11112_000011	(SLC5A2), mRNA
NM 000343	Homo sapiens solute carrier family 5 (sodium/glucose cotransporter), member 1
14147_000242	(SLC5A1), mRNA
NM_003040	Homo sapiens solute carrier family 4, anion exchanger, member 2 (erythrocyte
MM_003040	membrane protein band 3-like 1) (SLC4A2), mRNA
NR ( 000242	Homo sapiens solute carrier family 4, anion exchanger, member 1 (erythrocyte
NM_000342	membrane protein band 3, Diego blood group) (SLC4A1), mRNA
NR 6 000041	memorane protein baild 3, Diego blood group) (SEC4A1), micha
NM_000341	Homo sapiens solute carrier family 3 (cystine, dibasic and neutral amino acid
	transporters, activator of cystine, dibasic and neutral amino acid transport),
	member 1 (SLC3A1), mRNA
NM_001860	Homo sapiens solute carrier family 31 (copper transporters), member 2
	(SLC31A2), mRNA
NM_001859	Homo sapiens solute carrier family 31 (copper transporters), member 1
	(SLC31A1), mRNA
NM 003039	Homo sapiens solute carrier family 2 (facilitated glucose transporter), member 5
<del>-</del>	(SLC2A5), mRNA
NM_001042	Homo sapiens solute carrier family 2 (facilitated glucose transporter), member 4.
	(SLC2A4), mRNA
NM 003705	Homo sapiens solute carrier family 25 (mitochondrial carrier, Aralar), member
11112_005700	12 (SLC25A12), mRNA
NM_003060	Homo sapiens solute carrier family 22 (organic cation transporter), member 5
1111_005000	(SLC22A5), mRNA
NM_003058	Homo sapiens solute carrier family 22 (organic cation transporter), member 2
NM_003038	(SLC22A2), mRNA
NM 003057	Homo sapiens solute carrier family 22 (organic cation transporter), member 1
MM_003037	(SLC22A1), mRNA
NR 4 002560	Homo sapiens solute carrier family 25 (mitochondrial carrier; oxoglutarate
NM_003562	
	carrier), member 11 (SLC25A11), mRNA
NM_003038	Homo sapiens solute carrier family 1 (glutamate/neutral amino acid transporter),
	member 4 (SLC1A4), mRNA
NM_003056	Homo sapiens solute carrier family 19 (folate transporter), member 1
	(SLC19A1), mRNA
NM_003055	Homo sapiens solute carrier family 18 (vesicular acetylcholine), member 3
	(SLC18A3), mRNA
NM 003054	Homo sapiens solute carrier family 18 (vesicular monoamine), member 2
_	(SLC18A2), mRNA
NM 003053	Homo sapiens solute carrier family 18 (vesicular monoamine), member 1
	(SLC18A1), mRNA
NM_003052	Homo sapiens solute carrier family 34 (sodium phosphate), member 1
1111_005052	(SLC34A1), mRNA
NM 003051	Homo sapiens solute carrier family 16 (monocarboxylic acid transporters),
I COCOO TAINI	member 1 (SLC16A1), mRNA
NR 4 002004	Homo sapiens solute carrier family 13 (sodium-dependent dicarboxylate
NM_003984	
	transporter), member 2 (SLC13A2), mRNA

VM_000339	Homo sapiens solute carrier family 12 (sodium/chloride transporters), member 3
	(SLC12A3), mRNA Homo sapiens solute carrier family 12 (sodium/potassium/chloride transporters),
	member 2 (SLC12A2), mRNA Homo sapiens solute carrier family 10 (sodium/bile acid cotransporter family),
NM_003049	Homo sapiens solute carrier family 10 (sodium/bile acid cottansporter ransporter)
NM_003037	Homo sapiens signaling lymphocytic activation molecule (SLAM), mRNA Homo sapiens survival of motor neuron protein interacting protein 1 (SIP1),
NM_003616	Homo sapiens survival of motor neuron protein interacting protein
	mRNA Homo sapiens TAL1 (SCL) interrupting locus (SIL), mRNA  Homo sapiens TAL1 (SCL) interrupting locus (SIL), mRNA
NM_003035	Homo sapiens TALI (SCL) interrupting focus (SIL), interrupting focus (
NM_003032	(SIATI) mRNA
NM 001041	Home conjugate isomaltase (SI), mRNA
NM 003027	Transport CH3 domain (RR2-like 3 (SH3UL3), IIICVA
NM 003027	III comione SH3-domain GRB2-like 2 (SH3GL2), IIIKNA
NM 003025	TY Company GR B7-like [ (SH3ULT), IIIXIA
NM 003023	The strong strong the strong protein / (AD 1D) 41, 1110 MA
NM 003022	Homo sapiens SH3-domain binding protein 2 (SH3BGRL), Homo sapiens SH3 domain binding glutamic acid-rich protein like (SH3BGRL),
141410000022	
NM_000199	mRNA Homo sapiens N-sulfoglucosamine sulfohydrolase (sulfamidase) (SGSH),
NM_003020	mRNA Homo sapiens secretory granule, neuroendocrine protein 1 (7B2 protein)
_	
NM_000337	(SGNE1), mRNA  Homo sapiens sarcoglycan, delta (35kD dystrophin-associated glycoprotein)
	(SGCD), mRNA  Homo sapiens sarcoglycan, beta (43kD dystrophin-associated glycoprotein)
NM_000232	Homo sapiens sarcoglycan, beta (43kD dyshophin-associated gryo-pass)
	(SGCB), mRNA  Homo sapiens surfactant, pulmonary-associated protein D (SFTPD), mRNA
NM_003019	
NM_003018	
NM_000542	TOTAL CONTRACTOR ANY PROPERTY AND A STATE OF THE PROPERTY
NM_003011	Homo sapiens SET translocation (inycloid leakening)  Homo sapiens mitogen-activated protein kinase kinase 4 (MAP2K4), mRNA  Homo sapiens mitogen-activated protein kinase kinase 4 (MAP2K4), mRNA
NM_003010	Homo sapiens selenoprotein W, 1 (SEPW1), mRNA  Homo sapiens selenoprotein W, 1 (SEPW1), mRNA
NM_003009	Homo sapiens semenogelin II (SEMG2), mRNA
NM_003008	
NM_003007	
NM_003966	Homo sapiens sema domain, seven thromospondan repeats (springle like), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin)
NM_003002	and an enginete dehydrogenase complex, subulit D, littegral monitoring
14141_003002	(CDUD) nuclear gene encoding milocilondral protein, maxit
NM 002999	Homo sapiens syndecan 4 (amphiglycan, ryudocan) (SDC4); httq://
NM 002997	
NM 002996	Homo saniens small inducible cytokine subfamily D (Cys-A3-Cys), member 1
1111_002550	
NM 003175	Homo sapiens small inducible cytokine subtamily C, member 2 (80102),
1.2.2	1
NM 002993	Homo saniens small inducible cytokine subtamily B (Cys-A-Cys), memoer o
	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
NM_002994	inducible cytokine subtamily B (Cys-A-Cys), member s
	(epithelial-derived neutrophil-activating peptide 78) (SCYB5), mRNA

NM_002985	Homo sapiens small inducible cytokine A5 (RANTES) (SCYA5), mRNA
NM_002991	Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 24
_	(SCYA24), mRNA
NM 002990	Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 22
	(SCYA22), mRNA
NM 002989	Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 21
1111_002505	(SCYA21), mRNA
NM 002988	Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 18,
14141_002500	pulmonary and activation-regulated (SCYA18), mRNA
NM_002987	Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 17
14141_002507	(SCYA17), mRNA
NM_002986	Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 11
14141_002360	(eotaxin) (SCYA11), mRNA
NM 002979	Homo sapiens sterol carrier protein 2 (SCP2), mRNA
NM 001039	Homo sapiens sodium channel, nonvoltage-gated 1, gamma (SCNN1G), mRNA
	Homo sapiens sodium channel, nonvoltage-gated 1, delta (SCNN1D), mRNA
NM_002978	Homo sapiens sodium channel, nonvoltage-gated 1, detta (SCNN1D), mRNA
NM_001038	
NM_002977	Homo sapiens sodium channel, voltage-gated, type IX, alpha polypeptide
) To 6 000076	(SCN9A), mRNA
NM_002976	Homo sapiens sodium channel, voltage-gated, type VI, alpha polypeptide
77.6 000004	(SCN6A), mRNA
NM_000334	Homo sapiens sodium channel, voltage-gated, type IV, alpha polypeptide
27.5.004.005	(SCN4A), mRNA
NM_001037	Homo sapiens sodium channel, voltage-gated, type I, beta polypeptide (SCN1B),
12.00000	mRNA
NM_002975	Homo sapiens stem cell growth factor; lymphocyte secreted C-type lectin
77.5.000040	(SCGF), mRNA
NM_003843	Homo sapiens sciellin (SCEL), mRNA
NM_002973	Homo sapiens spinocerebellar ataxia 2 (olivopontocerebellar ataxia 2, autosomal
	dominant, ataxin 2) (SCA2), mRNA
NM_000332	Homo sapiens spinocerebellar ataxia 1 (olivopontocerebellar ataxia 1, autosomal
	dominant, ataxin 1) (SCA1), mRNA
NM_002971	Homo sapiens special AT-rich sequence binding protein 1 (binds to nuclear
	matrix/scaffold-associating DNA's) (SATB1), mRNA
NM_002970	Homo sapiens spermidine/spermine N1-acetyltransferase (SAT), mRNA
NM_003870	Homo sapiens IQ motif containing GTPase activating protein 1 (IQGAP1),
	mRNA
NM_002967	Homo sapiens scaffold attachment factor B (SAFB), mRNA
NM_000331	Homo sapiens serum amyloid A1 (SAA1), mRNA
NM_001036	Homo sapiens ryanodine receptor 3 (RYR3), mRNA
NM_001035	Homo sapiens ryanodine receptor 2 (cardiac) (RYR2), mRNA
NM_002956	Homo sapiens restin (Reed-Steinberg cell-expressed intermediate filament-
	associated protein) (RSN), mRNA
NM_001033	Homo sapiens ribonucleotide reductase M1 polypeptide (RRM1), mRNA
NM_002955	Homo sapiens ras responsive element binding protein 1 (RREB1), mRNA
NM_003942	Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 4 (RPS6KA4),
_	mRNA
NM_002953	Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 1 (RPS6KA1),
-	mRNA
NM 002951	Homo sapiens ribophorin II (RPN2), mRNA
NM 002950	Homo sapiens ribophorin I (RPN1), mRNA
NM 000329	Homo sapiens retinal pigment epithelium-specific protein (65kD) (RPE65),
	The state of the s

	mRNA (PD44) DVA
NM 002947	Homo sapiens replication protein A3 (14kD) (RPA3), mRNA
NM 002946	Homo saniens replication protein A2 (32kD) (RPA2), mRNA
NM 002945	Homo seniens replication protein A1 (70kD) (RPA1), mkNA
VM 000328	Homo sapiens retinitis pigmentosa GTPase regulator (RPGR), mRNA
NM 002943	Homo seniens RAR-related orphan receptor A (RORA), mRNA
NM 000327	Home saniens retinal outer segment membrane protein I (ROMI), mRNA
NM 003799	Homo sapiens RNA (guanine-7-) methyltransferase (RNMT), mRNA
NM_002939	Home ganiens ribonuclease/angiogenin inhibitor (RNH), mKNA
NM 003800	Homo sapiens RNA guanylyltransferase and 5'-phosphatase (RNGTT), mRNA
	Homo saniens ring finger protein 4 (RNF4), mRNA
NM_002938	Homo sapiens ATP-binding cassette, sub-family E (OABP), member 1
NM_002940	(ABCE1), mRNA
NTM 002026	Homo sapiens ribonuclease H1 (RNASEH1), mRNA
NM_002936	Homo sapiens ribonuclease, RNase A family, 3 (eosinophil cationic protein)
NM_002935	(RNASE3), mRNA
	Homo sapiens ribonuclease, RNase A family, 2 (liver, eosinophil-derived
NM_002934	Homo sapiens modificiences, Kivase ir mining, 2 (11-24, 11-24)
	neurotoxin) (RNASE2), mRNA  Homo sapiens RPB5-mediating protein (RMP), mRNA
NM_003796	Homo sapiens receptor-interacting serine-threonine kinase 2 (RIPK2), mRNA
NM_003821	Homo sapiens receptor-interacting serine-interacting serine-interactin
NM_003687_	Homo sapiens LIM domain protein (RIL), mRNA
NM_002929_	Homo sapiens rhodopsin kinase (RHOK), mRNA
NM_000324_	Homo sapiens Rhesus blood group-associated glycoprotein (RHAG), mRNA
NM_003835	Homo sapiens regulator of G-protein signalling 9 (RGS9), mRNA
NM_003617	Homo sapiens regulator of G-protein signalling 5 (RGS5), mRNA
NM 002923	Homo sapiens regulator of G-protein signalling 2, 24kD (RGS2), mRNA
NM 002922	Homo sapiens regulator of G-protein signalling 1 (RGS1), mRNA
NM 002928	Homo saniens regulator of G-protein signalling 16 (RGS16), mRNA
NM 002926	Homo sapiens regulator of G-protein signalling 12 (RGS12), mRNA
NM 003834	Homo sapiens regulator of G-protein signalling 11 (RGS11), mRNA
NM_002921	Homo saniens retinal G protein coupled receptor (RGR), mRNA
NM 000538	Homo saniens regulatory factor X-associated protein (RFXAP), mRNA
NM_003721	Homo sapiens regulatory factor X-associated ankyrin-containing protein
14141_005721	(DEVANK) mRNA
NM_002918	Homo sapiens regulatory factor X, 1 (influences HLA class II expression)
14141_002510	(PEY1) mRNA
NM 002916	Homo saniens replication factor C (activator 1) 4 (37kD) (RFC4), mRNA
NM 002915	Homo saniens replication factor C (activator 1) 3 (38kD) (RFC3), mRNA
NM 002914	Homo saniens replication factor C (activator 1) 2 (40kD) (RFC2), mRNA
	Homo sapiens gene with multiple splice variants near HD locus on 4p16.3
NM_003704	DECA 22) mDNA
ND4 002008	Homo sapiens v-rel avian reticuloendotheliosis viral oncogene homolog (REL),
NM_002908	mRNA
77.5 000000	Homo sapiens regenerating islet-derived 1 alpha (pancreatic stone protein,
NM_002909	pancreatic thread protein) (REG1A), mRNA
	Homo sapiens retinal degeneration, slow (retinitis pigmentosa 7) (RDS), mRNA
NM_000322	Homo sapiens retinal degeneration, slow (retinus pignientosa 7) (1000), mRNA  Homo sapiens retinal degeneration, slow (retinus pignientosa 7) (1000), mRNA
NM_002905	Homo sapiens retinoi denydrogenase 3 (11-cisand 3-cis) (12713), micros
NM_002903	Homo sapiens recoverin (RCV1), mRNA
NM_002902	Homo sapiens reticulocalbin 2, EF-hand calcium binding domain (RCN2),
	mRNA  The hand coloium binding domain (RCN1)
NM_002901	Homo sapiens reticulocalbin 1, EF-hand calcium binding domain (RCN1),
l .	mRNA

	DATA IN THE STATE A CORNAIN ASSESSMENT ASSES
NM_002896	Homo sapiens RNA binding motif protein 4 (RBM4), mRNA
NM_002895	Homo sapiens retinoblastoma-like 1 (p107) (RBL1), mRNA
NM_000321	Homo sapiens retinoblastoma 1 (including osteosarcoma) (RB1), mRNA
NM_000966	Homo sapiens retinoic acid receptor, gamma (RARG), mRNA
NM_000964	Homo sapiens retinoic acid receptor, alpha (RARA), mRNA
NM_002885	Homo sapiens RAP1, GTPase activating protein 1 (RAP1GA1), mRNA
NM_002884	Homo sapiens RAP1A, member of RAS oncogene family (RAP1A), mRNA
NM_002883	Homo sapiens Ran GTPase activating protein 1 (RANGAP1), mRNA
NM_002881	Homo sapiens v-ral simian leukemia viral oncogene homolog B (ras related; GTP binding protein) (RALB), mRNA
NM_002871	Homo sapiens RAB interacting factor (RABIF), mRNA
NM 003929	Homo sapiens RAB7, member RAS oncogene family-like 1 (RAB7L1), mRNA
NM 002869	Homo sapiens RAB6, member RAS oncogene family (RAB6), mRNA
NM 002868	Homo sapiens RAB5B, member RAS oncogene family (RAB5B), mRNA
NM 002867	Homo sapiens RAB3B, member RAS oncogene family (RAB3B), mRNA
NM 002866	Homo sapiens RAB3A, member RAS oncogene family (RAB3A), mRNA
NM 002870	Homo sapiens RAB13, member RAS oncogene family (RAB13), mRNA
NM 000320	Homo sapiens quinoid dihydropteridine reductase (QDPR), mRNA
NM 002864	Homo sapiens pregnancy-zone protein (PZP), mRNA
NM 002863	Homo sapiens phosphorylase, glycogen; liver (Hers disease, glycogen storage
	disease type VI) (PYGL), mRNA
NM 002862	Homo sapiens phosphorylase, glycogen; brain (PYGB), nuclear gene encoding
1411_002002	mitochondrial protein, mRNA
NM 002860	Homo sapiens pyrroline-5-carboxylate synthetase (glutamate gamma-
1411_002000	semialdehyde synthetase) (PYCS), mRNA
NM 000319	Homo sapiens peroxisome receptor 1 (PXR1), mRNA
NM 002859	Homo sapiens paxillin (PXN), mRNA
NM 002857	Homo sapiens peroxisomal farnesylated protein (PXF), mRNA
NM 002854	Homo sapiens parvalbumin (PVALB), mRNA
NM_002852	Homo sapiens pentaxin-related gene, rapidly induced by IL-1 beta (PTX3), mRNA
NM 000317	Homo sapiens 6-pyruvoyltetrahydropterin synthase (PTS), mRNA
NM 002851	Homo sapiens protein tyrosine phosphatase, receptor-type, Z polypeptide 1
11212_002001	(PTPRZ1), mRNA
NM 002850	Homo sapiens protein tyrosine phosphatase, receptor type, S (PTPRS), mRNA
NM 002846	Homo sapiens protein tyrosine phosphatase, receptor type, N (PTPRN), mRNA
NM 002845	Homo sapiens protein tyrosine phosphatase, receptor type, M (PTPRM), mRNA
NM 002844	Homo sapiens protein tyrosine phosphatase, receptor type, K (PTPRK), mRNA
NM 002843	Homo sapiens protein tyrosine phosphatase, receptor type, J (PTPRJ), mRNA
NM 002842	Homo sapiens protein tyrosine phosphatase, receptor type, H (PTPRH), mRNA
NM 002840	Homo sapiens protein tyrosine phosphatase, receptor type, F (PTPRF), mRNA
NM 002839	Homo sapiens protein tyrosine phosphatase, receptor type, D (PTPRD), mRNA
NM 002824	Homo sapiens parathymosin (PTMS), mRNA
NM 002823	Homo sapiens prothymosin, alpha (gene sequence 28) (PTMA), mRNA
NM 000316	Homo sapiens parathyroid hormone receptor 1 (PTHR1), mRNA
NM 002820	Homo sapiens parathyroid hormone-like hormone (PTHLH), mRNA
NM_000315	Homo sapiens parathyroid hormone (PTH), mRNA
NM_000960	Homo sapiens prostaglandin I2 (prostacyclin) receptor (IP) (PTGIR), mRNA
NM 000959	Homo sapiens prostaglandin E (prostacychii) receptor (II) (176IK), mRNA  Homo sapiens prostaglandin F receptor (FP) (PTGFR), mRNA
NM 000958	Homo sapiens prostaglandin F receptor (FF) (FTGFR), mRNA  Homo sapiens prostaglandin E receptor 4 (subtype EP4) (PTGER4), mRNA
NM 000957	Homo sapiens prostaglandin E receptor 4 (subtype EP4) (PTGER4), mRNA  Homo sapiens prostaglandin E receptor 3 (subtype EP3) (PTGER3), mRNA
NM 000955	Homo sapiens prostaglandin E receptor 1 (subtype EP1), 42kD (PTGER1),
1111 000333	1 Homo sapiens prostagianum E receptor 1 (Subtype Er 1), 42kD (1 10Ekt),

	mRNA
NM 000954	Homo saniens prostaglandin D2 synthase (21kD, brain) (PTGDS), mRNA
NM 000314	Homo sapiens phosphatase and tensin homolog (mutated in multiple advanced
_	cancers 1) (PTEN), mRNA
NM_000952	Home saniens platelet-activating factor receptor (PTAFR), mRNA
NM 002818	Homo sapiens proteasome (prosome, macropain) activator subunit 2 (PA28 beta)
	(PSME2) mRNA
NM_002811	Homo sapiens proteasome (prosome, macropain) 26S subunit, non-ATPase, 7
<u>-</u>	(Moy34 homolog) (PSMD7), mRNA
NM_002806	Homo sapiens proteasome (prosome, macropain) 26S subunit, ATPase, 6
14.12_002000	(PSMC6) mRNA
NM_002805	Homo sapiens proteasome (prosome, macropain) 26S subunit, ATPase, 5
1111_002000	(PSMC5) mRNA
NM_002804	Homo sapiens proteasome (prosome, macropain) 26S subunit, ATPase, 3
1111_00200	(PSMC3) mRNA
NM_002803	Homo sapiens proteasome (prosome, macropain) 26S subunit, ATPase, 2
1111_002000	(PSMC2) mRNA
NM_002802	Homo sapiens proteasome (prosome, macropain) 26S subunit, ATPase, 1
1111_002002	(PSMC1) mRNA
NM_002800	Homo sapiens proteasome (prosome, macropain) subunit, beta type, 9 (large
1414_002000	multifunctional protease 2) (PSMB9), mRNA
NM_002799	Homo sapiens proteasome (prosome, macropain) subunit, beta type, 7 (PSMB7),
14141_002755	mRNA
NM 002797	Homo sapiens proteasome (prosome, macropain) subunit, beta type, 5 (PSMB5),
14141_002777	mRNA
NM_002796	Homo sapiens proteasome (prosome, macropain) subunit, beta type, 4 (PSMB4),
1414_002750	mRNA
NM_002795	Homo sapiens proteasome (prosome, macropain) subunit, beta type, 3 (PSMB3),
1414_002/50	mRNA
NM_002794	Homo sapiens proteasome (prosome, macropain) subunit, beta type, 2 (PSMB2),
1111_002.5	mRNA
NM 002793	Homo sapiens proteasome (prosome, macropain) subunit, beta type, 1 (PSMB1),
1111_002.	mRNA
NM 002801	Homo sapiens proteasome (prosome, macropain) subunit, beta type, 10
14141_002001	(PSMB10), mRNA
NM 002790	Homo sapiens proteasome (prosome, macropain) subunit, alpha type, 5
1414_002750	(PSMA5), mRNA
NM_002788	Homo sapiens proteasome (prosome, macropain) subunit, alpha type, 3
14141_002700	(PSMA3), mRNA
NM 002786	Homo sapiens proteasome (prosome, macropain) subunit, alpha type, 1
14141_002700	(PSMA1), mRNA
NM 002783	Homo sapiens pregnancy specific beta-1-glycoprotein 7 (PSG7), mRNA
NM 002781	Homo sapiens pregnancy specific beta-1-glycoprotein 5 (PSG5), mRNA
NM 002781	Homo saniens pregnancy specific beta-1-glycoprotein 4 (PSG4), mRNA
	Homo sapiens pregnancy specific beta-1-glycoprotein 11 (Note redefinition of
NM_002785	symbol) (PSG11), mRNA
NM 002784	Homo sapiens pregnancy specific beta-1-glycoprotein 9 (PSG9), mRNA
NM 002784 NM 002779	Homo sapiens pleckstrin and Sec7 domain protein (PSD), mRNA
	Homo sapiens kallikrein 10 (KLK10), mRNA
NM_002776	Homo sapiens kallikrein 6 (neurosin, zyme) (KLK6), mRNA
NM_002774	Homo sapiens protease, serine, 8 (prostasin) (PRSS8), mRNA
NM_002773	
NM_002770	Homo sapiens protease, serme, 2 (4 years 2) (1 koo2), met 17

	(A) (DDGG1) DNA
NM_002769	Homo sapiens protease, serine, 1 (trypsin 1) (PRSS1), mRNA
NM_003619	Homo sapiens protease, serine, 12 (neurotrypsin, motopsin) (PRSS12), mRNA
NM_002775	Homo sapiens protease, serine, 11 (IGF binding) (PRSS11), mRNA
NM_002767	Homo sapiens phosphoribosyl pyrophosphate synthetase-associated protein 2 (PRPSAP2), mRNA
NM_002766	Homo sapiens phosphoribosyl pyrophosphate synthetase-associated protein 1 (PRPSAP1), mRNA
NM 002765	Homo sapiens phosphoribosyl pyrophosphate synthetase 2 (PRPS2), mRNA
NM 002764	Homo sapiens phosphoribosyl pyrophosphate synthetase 1 (PRPS1), mRNA
NM_003891	Homo sapiens protein Z, vitamin K-dependent plasma glycoprotein (PROZ), mRNA
NM 002763	Homo sapiens prospero-related homeobox 1 (PROX1), mRNA
NM 000313	Homo sapiens protein S (alpha) (PROS1), mRNA
NM_000312	Homo sapiens protein C (inactivator of coagulation factors Va and VIIIa) (PROC), mRNA
NM 002762	Homo sapiens protamine 2 (PRM2), mRNA
NM 002761	Homo sapiens protamine 1 (PRM1), mRNA
NM 000949	Homo sapiens prolactin receptor (PRLR), mRNA
NM 000948	Homo sapiens prolactin (PRL), mRNA
NM_002759	Homo sapiens protein kinase, interferon-inducible double stranded RNA dependent (PRKR), mRNA
NM 002756	Homo sapiens mitogen-activated protein kinase kinase 3 (MAP2K3), mRNA
NM 002749	Homo sapiens mitogen-activated protein kinase 7 (MAPK7), mRNA
NM 002745	Homo sapiens mitogen-activated protein kinase 1 (MAPK1), mRNA
NM 002751	Homo sapiens mitogen-activated protein kinase 11 (MAPK11), mRNA
NM 002753	Homo sapiens mitogen-activated protein kinase 10 (MAPK10), mRNA
NM_002743	Homo sapiens protein kinase C substrate 80K-H (PRKCSH), mRNA
NM_002742	Homo sapiens protein kinase C, mu (PRKCM), mRNA
NM_002741	Homo sapiens protein kinase C-like 1 (PRKCL1), mRNA
NM_002740	Homo sapiens protein kinase C, iota (PRKCI), mRNA
NM_002738	Homo sapiens protein kinase C, beta 1 (PRKCB1), mRNA
NM_002737	Homo sapiens protein kinase C, alpha (PRKCA), mRNA
NM_002736	Homo sapiens protein kinase, cAMP-dependent, regulatory, type II, beta (PRKAR2B), mRNA
NM_002734	Homo sapiens protein kinase, cAMP-dependent, regulatory, type I, alpha (tissue specific extinguisher 1) (PRKAR1A), mRNA
NM_002733	Homo sapiens protein kinase, AMP-activated, gamma 1 non-catalytic subunit (PRKAG1), mRNA
NM_002731	Homo sapiens protein kinase, cAMP-dependent, catalytic, beta (PRKACB), mRNA
NM_002730	Homo sapiens protein kinase, cAMP-dependent, catalytic, alpha (PRKACA), mRNA
NM 000947	Homo sapiens primase, polypeptide 2A (58kD) (PRIM2A), mRNA
NM 000946	Homo sapiens primase, polypeptide 1 (49kD) (PRIM1), mRNA
NM_002728	Homo sapiens proteoglycan 2, bone marrow (natural killer cell activator, eosinophil granule major basic protein) (PRG2), mRNA
NM 002727	Homo sapiens proteoglycan 1, secretory granule (PRG1), mRNA
NM 002726	Homo sapiens prolyl endopeptidase (PREP), mRNA
NM_002725	Homo sapiens proline arginine-rich end leucine-rich repeat protein (PRELP), mRNA
NM 002723	Homo sapiens proline-rich protein BstNI subfamily 4 (PRB4), mRNA
NM 002722	Homo sapiens pancreatic polypeptide (PPY), mRNA
	The second polypopular (2 = 7),

NM_000310	Homo sapiens palmitoyl-protein thioesterase 1 (ceroid-lipofuscinosis, neuronal 1,
	. 6 . 11 \ (T)TOTO 1 \
<b>–</b> ;	Homo sapiens protein phosphatase 4 (formerly X), catalytic subunit (PPP4C), mRNA
VM 002719	Homo sapiens protein phosphatase 2, regulatory subunit B (B56), gamma
	isoform (PPP2R5C), mRNA
	Homo sapiens protein phosphatase 2 (formerly 2A), catalytic subunit, alpha isoform (PPP2CA), mRNA
NM_002713	Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 8 (FFF 1K8),
7. 6 000710	mRNA Homo sapiens protein phosphatase 1, regulatory subunit 7 (PPP1R7), mRNA
NM_002712	Homo sapiens protein phosphatase 1, regulatory subunit 10 (PPP1R10), mRNA  Homo sapiens protein phosphatase 1, regulatory subunit 10 (PPP1R10), mRNA
NM_002714	Homo sapiens protein phosphatase 1, catalytic subunit, gamma isoform
NM_002710	(PRD1CC) DILA
NM_002709	Homo sapiens protein phosphatase 1, catalytic subunit, beta isoloriii (FFF 1CB),
	mRNA Homo sapiens protein phosphatase 1, catalytic subunit, alpha isoform (PPP1CA),
NM_002708	mRNA
NA 000200	Home conjens protoporphyringgen oxidase (PPOX), mRNA
NM_000309	Homo sapiens protein phosphatase 1B (formerly 2C), magnesium-dependent,
NM_002706	beta isoform (PPM1B), mRNA
NR 6 000705	Hame senions periplakin (PPI) mRNA
NM_002705	Home senions pentidylprolyl isomerase C (cyclophilin C) (PPIC), fikina
NM_000943	Homo sapiens protective protein for beta-galactosidase (galactosialidosis)
NM_000308	(DDCD) mDNA
37 f 000702	Homo sapiens phosphoribosyl pyrophosphate amidotransferase (PPAT), mRNA
NM_002703	Homo sapiens phosphatidic acid phosphatase type 2C (PPAP2C), mRNA
NM_003712	Homo sapiens phosphatidic acid phosphatase type 2B (PPAP2B), mRNA
NM_003713	Homo sapiens phosphatidic acid phosphatase type 2A (PPAP2A), mRNA  Homo sapiens phosphatidic acid phosphatase type 2A (PPAP2A), mRNA
NM_003711	Homo sapiens POU domain, class 6, transcription factor 1 (POU6F1), mRNA
NM_002702	Homo sapiens POU domain, class 5, transcription factor 1 (POU5F1), mRNA  Homo sapiens POU domain, class 5, transcription factor 1 (POU5F1), mRNA
NM_002701	Homo sapiens POU domain, class 3, transcription factor 3 (POU4F3), mRNA  Homo sapiens POU domain, class 4, transcription factor 3 (POU4F3), mRNA
NM_002700	Homo sapiens POU domain, class 4, transcription factor 4 (POU3F4), mRNA  Homo sapiens POU domain, class 3, transcription factor 4 (POU3F4), mRNA
NM_000307	Homo sapiens POU domain, class 3, transcription factor 1 (POU3F1), mRNA  Homo sapiens POU domain, class 3, transcription factor 1 (POU3F1), mRNA
NM_002699	Homo sapiens POU domain, class 3, transcription factor 1 (POU2F1), mRNA  Homo sapiens POU domain, class 2, transcription factor 1 (POU2F1), mRNA
NM_002697	Homo sapiens POU domain, class 2, transcription factor 1 (Pit1 growth hormon
NM_000306	Homo sapiens POU domain, class 1, transcription factor 1 (Pit1, growth hormone
	factor 1) (POU1F1), mRNA
NM_000446	Homo sapiens paraoxonase 1 (PON1), mRNA
NM_000939	Homo sapiens proopiomelanocortin (adrenocorticotropin/ beta-lipotropin/ alpha-
	melanocyte stimulating hormone/ beta-melanocyte stimulating hormone/ beta-
	endorphin) (POMC), mRNA
NM_002693	Homo sapiens polymerase (DNA directed), gamma (POLG), nuclear gene
	encoding mitochondrial protein, mRNA
NM_002692	Homo sapiens polymerase (DNA directed), epsilon 2 (POLE2), mRNA
NM_002691	Homo sapiens polymerase (DNA directed), delta 1, catalytic subunit (125kD) (POLD1), mRNA
NM 002690	Homo saniens polymerase (DNA directed), beta (POLB), mRNA
NM 002890 NM 003967	Homo saniens putative neurotransmitter receptor (PNR), mRNA
	Homo sapiens phenylethanolamine N-methyltransferase (PNMT), mRNA
NM_002686	Homo sapiens peripheral myelin protein 2 (PMP2), mRNA
THE COOKER	
NM_002677 NM_000304	Homo sapiens peripheral myelin protein 22 (PMP22), mRNA

NM 002674	Homo sapiens pro-melanin-concentrating hormone (PMCH), mRNA
NM 002668	Homo sapiens proteolipid protein 2 (colonic epithelium-enriched) (PLP2),
14141_002008	mRNA
NM_000935	Homo sapiens procollagen-lysine, 2-oxoglutarate 5-dioxygenase (lysine
14M_000733	hydroxylase) 2 (PLOD2), mRNA
NM 002667	Homo sapiens phospholamban (PLN), mRNA
NM 002666	Homo sapiens perilipin (PLIN), mRNA
NM 002665	Homo sapiens plasminogen-like (PLGL), mRNA
NM 000301	Homo sapiens plasminogen (PLG), mRNA
NM 000445	Homo sapiens plectin 1, intermediate filament binding protein, 500kD (PLEC1),
14141_000443	mRNA
NM 002663	Homo sapiens phospholipase D2 (PLD2), mRNA
NM 002662	Homo sapiens phospholipase D1, phophatidylcholine-specific (PLD1), mRNA
NM_002661	Homo sapiens phospholipase C, gamma 2 (phosphatidylinositol-specific)
14M_002001	(PLCG2), mRNA
NM_002660	Homo sapiens phospholipase C, gamma 1 (formerly subtype 148) (PLCG1),
14141_002000	mRNA
NM 000933	Homo sapiens phospholipase C, beta 4 (PLCB4), mRNA
NM 002659	Homo sapiens plasminogen activator, urokinase receptor (PLAUR), mRNA
NM 002658	Homo sapiens plasminogen activator, urokinase (PLAU), mRNA
	Homo sapiens pleiomorphic adenoma gene 1 (PLAG1), mRNA
NM_002655	Homo sapiens phesoniolipine adenoma gene i (i Erker), macking the Homo sapiens phospholipase A2, group V (PLA2G5), mRNA
NM_000929	Homo sapiens phospholipase A2, group V (12/2007), interview Homo sapiens phospholipase A2, group IVC (cytosolic, calcium-independent)
NM_003706	(PLA2G4C), mRNA
NM 000300	Homo sapiens phospholipase A2, group IIA (platelets, synovial fluid)
1/1/1 _000300	(PLA2G2A), nuclear gene encoding mitochondrial protein, mRNA
NM 003561	Homo sapiens phospholipase A2, group X (PLA2G10), mRNA
NM 002654	Homo sapiens pyruvate kinase, muscle (PKM2), mRNA
	Homo sapiens serine/threonine kinase 16 (STK16), mRNA
NM_003691	Homo sapiens polycystic kidney disease 1 (autosomal dominant) (PKD1),
NM_000296	mRNA
NM_003607	Homo sapiens Ser-Thr protein kinase related to the myotonic dystrophy protein
1003007	kinase (PK428), mRNA
NM 003678	Homo sapiens gene from NF2/meningioma region of 22q12 (PK1.3), mRNA
	Homo sapiens paired-like homeodomain transcription factor 2 (PITX2), mRNA
NM_000325	Homo sapiens paired-like homeodomain transcription factor 1 (PITX1), mRNA
NM_002653	Homo sapiens prolactin-induced protein (PIP), mRNA
NM_002652	Homo sapiens protactin-induced protein (FIF), individual Homo sapiens phosphatidylinositol-4-phosphate 5-kinase, type I, beta
NM_003558	
ND 4 002555	(PIP5K1B), mRNA  Homo sapiens phosphatidylinositol-4-phosphate 5-kinase, type I, alpha
NM_003557	
ND 6 002746	(PIP5K1A), mRNA  Homo sapiens dynein, cytoplasmic, light polypeptide (PIN), mRNA
NM_003746	
NM_002648	Homo sapiens pim-1 oncogene (PIM1), mRNA  Homo sapiens phosphatidylinositol 4-kinase, catalytic, beta polypeptide
NM_002651	
ND 6 0000610	(PIK4CB), mRNA
NM_002643	Homo sapiens phosphatidylinositol glycan, class F (PIGF), mRNA
NM_002642	Homo sapiens phosphatidylinositol glycan, class C (PIGC), mRNA
NM_002638	Homo sapiens protease inhibitor 3, skin-derived (SKALP) (PI3), mRNA
NM_000294	Homo sapiens phosphorylase kinase, gamma 2 (testis) (PHKG2), mRNA
NM_000293	Homo sapiens phosphorylase kinase, beta (PHKB), mRNA
NM_000292	Homo sapiens phosphorylase kinase, alpha 2 (liver) (PHKA2), mRNA
NM_002637	Homo sapiens phosphorylase kinase, alpha 1 (muscle) (PHKA1), mRNA

	TY (DCD) mDNA
NM_000926	Homo sapiens progesterone receptor (PGR), mRNA
NM_002633	Homo sapiens phosphoglucomutase 1 (PGM1), mRNA
NM_000291	Homo sapiens phosphoglycerate kinase 1 (PGK1), mRNA  Homo sapiens placental growth factor, vascular endothelial growth factor-related
NM_002632	Homo sapiens placental growth factor, vascular endomenal growth factor-related
	protein (PGF), mRNA
NM_002631	Homo sapiens phosphogluconate dehydrogenase (PGD), mRNA
NM_002630	Homo sapiens progastricsin (pepsinogen C) (PGC), mRNA
NM_000290	Homo sapiens phosphoglycerate mutase 2 (muscle) (PGAM2), mRNA
NM_002629	Homo sapiens phosphoglycerate mutase 1 (brain) (PGAM1), mRNA  Homo sapiens phosphofructokinase, muscle (PFKM), mRNA
NM_000289	Homo sapiens phosphofructokinase, inuscie (FKKL), mRNA
NM_002626	Homo sapiens 6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 1
NM_002625	(PFKFB1), mRNA
NM 002621	Homo sapiens properdin P factor, complement (PFC), mRNA
	Homo sapiens platelet factor 4 variant 1 (PF4V1), mRNA
NM 002620	Homo sapiens platelet factor 4 (PF4), mRNA
NM_002619	Homo sapiens peroxisomal biogenesis factor 7 (PEX7), mRNA
NM_000288 NM_000287	Homo sapiens peroxisomal biogenesis factor 6 (PEX6), mRNA
NM 003630	Homo sapiens peroxisomal biogenesis factor 3 (PEX3), mRNA
NM 000466	Homo sapiens peroxisome biogenesis factor 1 (PEX1), mRNA
NM 002618	Homo sapiens peroxisome biogenesis factor 13 (PEX13), mRNA
NM 000442	Homo sapiens platelet/endothelial cell adhesion molecule (CD31 antigen)
NWI_000442	(PECAM1), mRNA
NM 002614	Homo sapiens PDZ domain containing 1 (PDZK1), mRNA
NM_003477	Homo sapiens Pyruvate dehydrogenase complex, lipoyl-containing component
14141_005477	X; E3-binding protein (PDX1), mRNA
NM 002613	Homo sapiens 3-phosphoinositide dependent protein kinase-1 (PDPK1), mRNA
NM 002612	Homo sapiens pyruvate dehydrogenase kinase, isoenzyme 4 (PDK4), mRNA
NM 000925	Homo sapiens pyruvate dehydrogenase (lipoamide) beta (PDHB), mRNA
NM 000284	Homo sapiens pyruvate dehydrogenase (lipoamide) alpha 1 (PDHA1), mRNA
NM 000924	Homo sapiens phosphodiesterase IB, calmodulin-dependent (PDE1B), mRNA
NM 002606	Homo sapiens phosphodiesterase 9A (PDE9A), mRNA
NM 002602	Homo sapiens phosphodiesterase 6G, cGMP-specific, rod, gamma (PDE6G),
	mRNA
NM_002601	Homo sapiens phosphodiesterase 6D, cGMP-specific, rod, delta (PDE6D),
	mRNA
NM_000921	Homo sapiens phosphodiesterase 3A, cGMP-inhibited (PDE3A), mRNA
NM_002598	Homo sapiens programmed cell death 2 (PDCD2), mRNA
NM_002594	Homo sapiens proprotein convertase subtilisin/kexin type 2 (PCSK2), mRNA
NM_002592	Homo sapiens proliferating cell nuclear antigen (PCNA), mRNA
NM_002591	Homo sapiens phosphoenolpyruvate carboxykinase 1 (soluble) (PCK1), mRNA
NM_002586	Homo sapiens pre-B-cell leukemia transcription factor 2 (PBX2), mRNA
NM_002585	Homo sapiens pre-B-cell leukemia transcription factor 1 (PBX1), mRNA
NM_002583	Homo sapiens PRKC, apoptosis, WT1, regulator (PAWR), mRNA
NM_002582	Homo sapiens poly(A)-specific ribonuclease (deadenylation nuclease) (PARN), mRNA
NM 003631	Homo sapiens poly (ADP-ribose) glycohydrolase (PARG), mRNA
NM 002580	Homo sapiens pancreatitis-associated protein (PAP), mRNA
NM_000919	Homo sapiens peptidylglycine alpha-amidating monooxygenase (PAM), mRNA
	Homo sapiens p21 (CDKN1A)-activated kinase 3 (PAK3), mRNA
NM_002578	
NM_002574	Homo sapiens peroxiredoxin 1 (PRDX1), mRNA  Homo sapiens platelet-activating factor acetylhydrolase, isoform lb, gamma

nit (29kD) (PAFAH1B3), mRNA
o sapiens platelet-activating factor acetylhydrolase, isoform Ib, beta subunit () (PAFAH1B2), mRNA
sapiens progestagen-associated endometrial protein (placental protein 14,
ancy-associated endometrial alpha-2-globulin, alpha uterine protein)
P), mRNA
o sapiens paired basic amino acid cleaving enzyme (furin, membrane
iated receptor protein) (PACE), mRNA
o sapiens paired basic amino acid cleaving system 4 (PACE4), mRNA
o sapiens sequestosome 1 (SQSTM1), mRNA
o sapiens procollagen-proline, 2-oxoglutarate 4-dioxygenase (proline 4-
oxylase), beta polypeptide (protein disulfide isomerase; thyroid hormone
ng protein p55) (P4HB), mRNA
o sapiens procollagen-proline, 2-oxoglutarate 4-dioxygenase (proline 4-
oxylase), alpha polypeptide I (P4HA1), mRNA
o sapiens pyrimidinergic receptor P2Y, G-protein coupled, 4 (P2RY4),
[A
o sapiens purinergic receptor P2Y, G-protein coupled, 2 (P2RY2), mRNA
o sapiens purinergic receptor P2Y, G-protein coupled, 11 (P2RY11),
IA
o sapiens purinergic receptor P2X, ligand-gated ion channel, 7 (P2RX7),
JA
o sapiens purinergic receptor P2X, ligand-gated ion channel, 5 (P2RX5),
VA
o sapiens purinergic receptor P2X, ligand-gated ion channel, 4 (P2RX4),
NA
to sapiens purinergic receptor P2X, ligand-gated ion channel, 3 (P2RX3),
NA
no sapiens oxysterol binding protein (OSBP), mRNA
no sapiens orosomucoid 2 (ORM2), mRNA
no sapiens olfactory receptor, family 6, subfamily A, member 1 (OR6A1),
NA
no sapiens olfactory receptor, family 3, subfamily A, member 1 (OR3A1),
NA
no sapiens olfactory receptor, family 1, subfamily D, member 2 (OR1D2),
NA
no sapiens opioid receptor, mu 1 (OPRM1), mRNA
no sapiens opioid receptor, kappa 1 (OPRK1), mRNA
no sapiens opioid receptor, delta 1 (OPRD1), mRNA
no sapiens oligodendrocyte myelin glycoprotein (OMG), mRNA
no sapiens oxidised low density lipoprotein (lectin-like) receptor 1 (OLR1),
NA
no sapiens G protein-coupled receptor 68 (GPR68), mRNA
no sapiens outer dense fibre of sperm tails 2 (ODF2), mRNA
no sapiens nuclear VCP-like (NVL), mRNA
no sapiens neurotensin receptor 1 (high affinity) (NTSR1), mRNA
no sapiens neurotrophic tyrosine kinase, receptor, type 3 (NTRK3), mRNA
no sapiens 5' nucleotidase (CD73) (NT5), mRNA
no sapiens neutral sphingomyelinase (N-SMase) activation associated factor
MAF), mRNA
,, · · - · ·
no sapiens ectodermal-neural cortex (with BTB-like domain) (ENC1),

NM 003872	Homo sapiens neuropilin 2 (NRP2), mRNA
NM 003873	Homo sapiens neuropilin 1 (NRP1), mRNA
NM 003489	Homo sapiens nuclear receptor interacting protein 1 (NRIP1), mRNA
NM 002525	Homo sapiens nardilysin (N-arginine dibasic convertase) (NRD1), hikkya
NM 000905	Homo saniens neuropeptide Y (NPY), mRNA
NM 000910	Homo saniens neuropeptide Y receptor Y2 (NPY2R), mRNA
NM 000909	Homo sapiens neuropeptide Y receptor YI (NPYIK), mRNA
NM 002522	t = = = = = = = 1 = ontrovin I (NPTX I) mRIVA
NM_000908	Homo sapiens natriuretic peptide receptor C/guanylate cyclase C (atrionatiumetic
NM_000906	Homo sapiens natriuretic peptide receptor A/guanylate cyclase A (atrionatriuretic peptide receptor A) (NPR1), mRNA
	Homo sapiens natriuretic peptide precursor B (NPPB), mRNA
NM_002521	Homo sapiens nuclear protein, ataxia-telangiectasia locus (NPAT), mRNA
NM_002519	Homo sapiens nuclear protein, ataxia-telangicousta 1993 (NPAS2) mRNA
NM_002518	Homo sapiens neuronal PAS domain protein 2 (NPAS2), mRNA  Homo sapiens neuronal PAS domain protein 1 (NPAS1), mRNA
NM_002517	Homo sapiens neuronal PAS domain protein 1 (NPAS1), mRNA  Homo sapiens neuronal PAS domain protein 1 (NPAS1), mRNA
NM_002514	Homo sapiens nephroblastoma overexpressed gene (NOV), mRNA
NM 003787	Homo sapiens nucleolar protein 4 (NOL4), mRNA
NM_003946	Homo sapiens nucleolar protein 3 (apoptosis repressor with CARD domain)
NM_003551	Homo sapiens non-metastatic cells 5, protein expressed in (nucleoside-
11112_000000	diphographyte kingse) (NME5) mRNA
NM 002513	Transa capione non-metastatic cells 3 protein expressed in (NIVIES), IIINIA
NM_002512	Homo saniens non-metastatic cells 2, protein (NM23B) expressed in (NM22),
NW_002312	nuclear gene encoding mitochondrial protein, mRNA
NM 002511	Home series regromed in R recentor (NMBR), mRNA
	Transported (transmembrane) nmb (GPNMB), IIINNA
NM_002510	Homo sapiens mitogen-activated protein kinase kinase kinase 14 (MAP3K14),
NM_003954	mRNA
37 6 000500	TY regions nidogen (enactin) (NID) mRNA
NM_002508	Homo sapiens nerve growth factor receptor (TNFR superfamily, member 16)
NM_002507	Homo sapiens nerve growth factor recopier (111224 by
	(NGFR), mRNA  Homo sapiens nerve growth factor, beta polypeptide (NGFB), mRNA
NM_002506	Homo sapiens nerve growth factor, octa polypeptide (1022); Homo sapiens nuclear factor of kappa light polypeptide gene enhancer in B-cells
NM_002503	1: 1: 1-item hata (NEKRIR) mRNA
NM_002502	Homo sapiens nuclear factor of kappa light polypeptide gene enhancer in B-cells 2 (p49/p100) (NFKB2), mRNA
NM_002501	Homo sapiens nuclear factor I/X (CCAAT-binding transcription factor) (NFIX), mRNA
NM 002500	TY comions neurogenic differentiation 1 (NEUROD1), mRNA
NM 002497	Homo saniens NIMA (never in mitosis gene a)-related kinase 2 (NEX2), history
NM_002496	Homo sapiens NADH dehydrogenase (ubiquinone) Fe-S protein 8 (25kD)
	(NADH-coenzyme Q reductase) (NDUFS8), mRNA
NM_002495	OIADU goenzame O reductase) (NI)UES4), MKNA
NM_002494	Homo sapiens NADH dehydrogenase (ubiquinone) 1, subcomplex unknown, 1
NM_002490	Homo sapiens NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 6
NM_002488	Homo sapiens NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 2 (8kL
27 2 22 22 2	Homo ganiens N. descetules a N. sulfotransferase (henaran glucosaminyl) 2
NM_002490	(6kD, KFYI) (NDUFC1), mRNA  Homo sapiens NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 6 (14kD, B14) (NDUFA6), mRNA  Homo sapiens NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 2 (8kB8) (NDUFA2), mRNA

	(NDST2), mRNA
NM 001543	Homo sapiens N-deacetylase/N-sulfotransferase (heparan glucosaminyl) 1
1111_001545	(NDST1), mRNA
NM 003581	Homo sapiens NCK adaptor protein 2 (NCK2), mRNA
NM 002486	Homo sapiens nuclear cap binding protein subunit 1, 80kD (NCBP1), mRNA
NM 002483	Homo sapiens carcinoembryonic antigen-related cell adhesion molecule 6 (non-
1111_002.00	specific cross reacting antigen) (CEACAM6), mRNA
NM_000662	Homo sapiens N-acetyltransferase 1 (arylamine N-acetyltransferase) (NAT1),
_	mRNA
NM_000263	Homo sapiens N-acetylglucosaminidase, alpha- (Sanfilippo disease IIIB)
	(NAGLÚ), mRNA
NM 003871	Homo sapiens myelin transcription factor 2 (MYT2), mRNA
NM 003803	Homo sapiens myomesin 1 (skelemin) (185kD) (MYOM1), mRNA
NM 002479	Homo sapiens myogenin (myogenic factor 4) (MYOG), mRNA
NM 002472	Homo sapiens myosin, heavy polypeptide 8, skeletal muscle, perinatal (MYH8),
	mRNA
NM_002469	Homo sapiens myogenic factor 6 (herculin) (MYF6), mRNA
NM_002468	Homo sapiens myeloid differentiation primary response gene (88) (MYD88),
	mRNA
NM_002460	Homo sapiens interferon regulatory factor 4 (IRF4), mRNA
NM_002457	Homo sapiens mucin 2, intestinal/tracheal (MUC2), mRNA
NM_002456	Homo sapiens mucin 1, transmembrane (MUC1), mRNA
NM_002455	Homo sapiens metaxin 1 (MTX1), mRNA
NM_002453	Homo sapiens mitochondrial translational initiation factor 2 (MTIF2), nuclear
	gene encoding mitochondrial protein, mRNA
NM_002452	Homo sapiens nudix (nucleoside diphosphate linked moiety X)-type motif 1
ND 6 000 450	(NUDT1), mRNA
NM_002450	Homo sapiens metallothionein 1L (MT1L), mRNA
NM_002447	Homo sapiens macrophage stimulating 1 receptor (c-met-related tyrosine kinase) (MST1R), mRNA
NM_002446	Homo sapiens mitogen-activated protein kinase kinase kinase 10 (MAP3K10),
	mRNA
NM_002445	Homo sapiens macrophage scavenger receptor 1 (MSR1), mRNA
NM_002444	Homo sapiens moesin (MSN), mRNA
NM_003879	Homo sapiens CASP8 and FADD-like apoptosis regulator (CFLAR), mRNA
NM_000530	Homo sapiens myelin protein zero (Charcot-Marie-Tooth neuropathy 1B) (MPZ), mRNA
NM_002437	Homo sapiens MpV17 transgene, murine homolog, glomerulosclerosis (MPV17), mRNA
NM_001932	Homo sapiens membrane protein, palmitoylated 3 (MAGUK p55 subfamily member 3) (MPP3), mRNA
NM 002435	Homo sapiens mannose phosphate isomerase (MPI), mRNA
NM_002433	Homo sapiens N-methylpurine-DNA glycosylase (MPG), mRNA
NM 003829	Homo sapiens N-methylputhie-DNA glycosylase (MPG), mRNA  Homo sapiens multiple PDZ domain protein (MPDZ), mRNA
NM 003829	Homo sapiens Fas (TNFRSF6)-associated via death domain (FADD), mRNA
NM 002432	Homo sapiens myeloid cell nuclear differentiation antigen (MNDA), mRNA
NM 002432	Homo sapiens menage a trois 1 (CAK assembly factor) (MNAT1), mRNA
NM 002431	Homo sapiens meningioma (disrupted in balanced translocation) 1 (MN1),
	mRNA
NM_000901	Homo sapiens nuclear receptor subfamily 3, group C, member 2 (NR3C2), mRNA
NM 003482	Homo sapiens myeloid/lymphoid or mixed-lineage leukemia 2 (MLL2), mRNA

	11 (0.54 DOTZ11)
NM_002419	Homo sapiens mitogen-activated protein kinase kinase kinase 11 (MAP3K11),
	l mRNA
NM_002417	Homo sapiens antigen identified by monoclonal antibody Ki-67 (MKI67),
	mRNA
NM_002416	Homo sapiens monokine induced by gamma interferon (MIG), mRNA
NM_002415	Homo sapiens macrophage migration inhibitory factor (glycosylation-inhibiting
_	factor) (MIF), mRNA
NM 002413	Homo sapiens microsomal glutathione S-transferase 2 (MGST2), mRNA
NM 000900	Homo sapiens matrix Gla protein (MGP), mRNA
NM 002412	Homo sapiens O-6-methylguanine-DNA methyltransferase (MGMT), mRNA
NM 002407	Homo sapiens mammaglobin 2 (MGB2), mRNA
NM 002411	Homo sapiens mammaglobin 1 (MGB1), mRNA
NM_002397	Homo sapiens MADS box transcription enhancer factor 2, polypeptide C
14141_002557	(myocyte enhancer factor 2C) (MEF2C), mRNA
NM 002391	Homo sapiens midkine (neurite growth-promoting factor 2) (MDK), mRNA
NM 002387	Homo sapiens mutated in colorectal cancers (MCC), mRNA
NM_000529	Homo sapiens melanocortin 2 receptor (adrenocorticotropic hormone) (MC2R),
14141_000525	mRNA
NM_002386	Homo sapiens melanocortin 1 receptor (alpha melanocyte stimulating hormone
NWI_002380	receptor) (MC1R), mRNA
NM 002385	Homo sapiens myelin basic protein (MBP), mRNA
NM 002383	Homo sapiens MAX protein (MAX), mRNA
	Homo sapiens megakaryocyte-associated tyrosine kinase (MATK), mRNA
NM_002378	Homo sapiens MAP/microtubule affinity-regulating kinase 3 (MARK3), mRNA
NM_002376	Homo sapiens monoamine oxidase B (MAOB), nuclear gene encoding
NM_000898	Homo sapiens monoanine oxidase B (WAOB), nuclear gene choosing
ND 4 002 400	mitochondrial protein, mRNA  Homo sapiens Microfibril-associated glycoprotein-2 (MAGP2), mRNA
NM_003480	Homo sapiens micronomi-associated grycoprotein-2 (MAGEB4), mRNA  Homo sapiens melanoma antigen, family B, 4 (MAGEB4), mRNA
NM_002367	Homo sapiens melanoma antigen, family B, 3 (MAGEB3), mRNA  Homo sapiens melanoma antigen, family B, 3 (MAGEB3), mRNA
NM_002365	Homo sapiens melanoma antigen, family B, 3 (MAGEB3), mRNA  Homo sapiens melanoma antigen, family B, 2 (MAGEB2), mRNA
NM_002364	Homo sapiens melanoma antigen, family B, 2 (WAGED2), midvix
NM_002363	Homo sapiens melanoma antigen, family B, 1 (MAGEB1), mRNA
NM_002362	Homo sapiens melanoma antigen, family A, 4 (MAGEA4), mRNA
NM_003682	Homo sapiens MAP-kinase activating death domain (MADD), mRNA
NM_002357	Homo sapiens MAX dimerization protein (MAD), mRNA
NM_002350	Homo sapiens v-yes-1 Yamaguchi sarcoma viral related oncogene homolog
	(LYN), mRNA
NM_002349	Homo sapiens lymphocyte antigen 75 (LY75), mRNA
NM_002347	Homo sapiens lymphocyte antigen 6 complex, locus H (LY6H), mRNA
NM_002346	Homo sapiens lymphocyte antigen 6 complex, locus E (LY6E), mRNA
NM_002345	Homo sapiens lumican (LUM), mRNA
NM_002344	Homo sapiens leukocyte tyrosine kinase (LTK), mRNA
NM_002343	Homo sapiens lactotransferrin (LTF), mRNA
NM_000897	Homo sapiens leukotriene C4 synthase (LTC4S), mRNA
NM_003573	Homo sapiens latent transforming growth factor beta binding protein 4 (LTBP4),
	mRNA
NM_000752	Homo sapiens leukotriene b4 receptor (chemokine receptor-like 1) (LTB4R),
_	mRNA
NM 000895	Homo sapiens leukotriene A4 hydrolase (LTA4H), mRNA
NM 002340	Homo sapiens lanosterol synthase (2,3-oxidosqualene-lanosterol cyclase) (LSS),
	mRNA
NM 002338	Homo sapiens limbic system-associated membrane protein (LSAMP), mRNA
NM 002337	Homo sapiens low density lipoprotein-related protein-associated protein 1

	(alpha-2-macroglobulin receptor-associated protein 1) (LRPAP1), mRNA
NM 002336	Homo sapiens low density lipoprotein receptor-related protein 6 (LRP6), mRNA
NM 002330	Homo sapiens leucine-rich neuronal protein (LRN), mRNA
NM 002317	Homo sapiens lysyl oxidase (LOX), mRNA
NM 002317	Homo sapiens LIM homeobox transcription factor 1, beta (LMX1B), mRNA
NM 002315	Homo sapiens LIM domain only 1 (rhombotin 1) (LMO1), mRNA
NM 002313	Homo sapiens ligase IV, DNA, ATP-dependent (LIG4), mRNA
NM 002312	Homo sapiens lectin, galactoside-binding, soluble, 3 (galectin 3) (LGALS3),
1414_002300	mRNA
NM 002303	Homo sapiens leptin receptor (LEPR), mRNA
NM 002302	Homo sapiens leukocyte cell-derived chemotaxin 2 (LECT2), mRNA
NM 001290	Homo sapiens LIM domain binding 2 (LDB2), mRNA
NM 003893	Homo sapiens LIM domain binding 1 (LDB1), mRNA
NM 002299	Homo sapiens lactase (LCT), mRNA
NM 002297	Homo sapiens lipocalin 1 (protein migrating faster than albumin, tear
_	prealbumin) (LCN1), mRNA
NM_002296	Homo sapiens lamin B receptor (LBR), mRNA
NM_002291	Homo sapiens laminin, beta 1 (LAMB1), mRNA
NM 002289	Homo sapiens lactalbumin, alpha- (LALBA), mRNA
NM 002273	Homo sapiens keratin 8 (KRT8), mRNA
NM 002276	Homo sapiens keratin 19 (KRT19), mRNA
NM 002275	Homo sapiens keratin 15 (KRT15), mRNA
NM 002274	Homo sapiens keratin 13 (KRT13), mRNA
NM 002265	Homo sapiens karyopherin (importin) beta 1 (KPNB1), mRNA
NM 002267	Homo sapiens karyopherin alpha 3 (importin alpha 4) (KPNA3), mRNA
NM_002266	Homo sapiens karyopherin alpha 2 (RAG cohort 1, importin alpha 1) (KPNA2), mRNA
NM 000893	Homo sapiens kininogen (KNG), mRNA
NM 003679	Homo sapiens kynurenine 3-monooxygenase (kynurenine 3-hydroxylase)
1414_005075	(KMO), mRNA
NM_002258	Homo sapiens killer cell lectin-like receptor subfamily B, member 1 (KLRB1), mRNA
NM 002257	Homo sapiens kallikrein 1, renal/pancreas/salivary (KLK1), mRNA
NM 002256	Homo sapiens KiSS-1 metastasis-suppressor (KISS1), mRNA
NM 002255	Homo sapiens killer cell immunoglobulin-like receptor, two domains, long
14141_002255	cytoplasmic tail, 4 (KIR2DL4), mRNA
NM 002254	Homo sapiens kinesin family member 3C (KIF3C), mRNA
NM 003958	Homo sapiens ring finger protein (C3HC4 type) 8 (RNF8), mRNA
NM 003685	Homo sapiens KH-type splicing regulatory protein (FUSE binding protein 2)
1111_005005	(KHSRP), mRNA
NM 002252	Homo sapiens potassium voltage-gated channel, delayed-rectifier, subfamily S,
1111_002202	member 3 (KCNS3), mRNA
NM 002250	Homo sapiens potassium intermediate/small conductance calcium-activated
1111_002200	channel, subfamily N, member 4 (KCNN4), mRNA
NM_002249	Homo sapiens potassium intermediate/small conductance calcium-activated
	channel, subfamily N, member 3 (KCNN3), mRNA
NM 002247	Homo sapiens potassium large conductance calcium-activated channel,
	subfamily M, alpha member 1 (KCNMA1), mRNA
NM 002244	Homo sapiens potassium inwardly-rectifying channel, sublamily J, inhibitor I
NM_002244	Homo sapiens potassium inwardly-rectifying channel, subfamily J, inhibitor 1 (KCNJN1), mRNA
NM_002244 NM_002240	

NM_002239	Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 3 (KCNJ3), mRNA
NM_000891	Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 2
	(KCNJ2), mRNA
NM_002241	Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 10 (KCNJ10), mRNA
NM_002238	Homo sapiens potassium voltage-gated channel, subfamily H (eag-related),
NM_002237	Homo sapiens potassium voltage-gated channel, subfamily G, member 1
NM_002236	Homo sapiens potassium voltage-gated channel, subfamily F, member 1
NM 003636	Homo sapiens potassium voltage-gated channel, shaker-related subfamily, beta
	member 2 (KCNAR2) mRNA
NM_003471	Homo sapiens potassium voltage-gated channel, shaker-related subfamily, beta member 1 (KCNAB1), mRNA
NM_002235	Homo sapiens potassium voltage-gated channel, shaker-related subfamily,
1,1,1002200	member 6 (KCNA6) mRNA
NM_002234	Homo sapiens potassium voltage-gated channel, shaker-related subfamily,
NM_002233	Homo sapiens potassium voltage-gated channel, shaker-related subfamily, member 4 (KCNA4), mRNA
NM 002232	Homo sapiens potassium voltage-gated channel, shaker-related subfamily,
<del></del>	member 3 (KCNA3), mRNA
NM 002229	Homo seniens in B proto-oncogene (IUNB), mRNA
NM 003666	Homo saniens hasic leucine zipper nuclear factor 1 (JEM-1) (BLZF1), mRNA
NM 002227	Homo sapiens Janus kinase 1 (a protein tyrosine kinase) (JAK1), mRNA
NM 003024	Homo saniens intersectin 1 (SH3 domain protein) (ITSN1), mRNA
NM 002224	Homo sapiens inositol 1,4,5-triphosphate receptor, type 3 (ITPR3), mRNA
NM 002223	Homo sapiens inositol 1,4,5-triphosphate receptor, type 2 (ITPR2), mRNA
NM 002221	Homo sapiens inositol 1,4,5-trisphosphate 3-kinase B (ITPKB), mRNA
NM 002220	Homo sapiens inositol 1,4,5-trisphosphate 3-kinase A (ITPKA), mRNA
NM 002220	Homo sapiens integral membrane protein 1 (ITM1), mRNA
NM_002219	Homo sapiens inter-alpha (globulin) inhibitor H4 (plasma Kallıkrein-sensitive
37.6.000016	glycoprotein) (ITIH4), mRNA  Homo sapiens inter-alpha (globulin) inhibitor, H2 polypeptide (ITIH2), mRNA
NM_002216	Homo sapiens inter-alpha (globulin) inhibitor, H1 polypeptide (ITIH1), mRNA
NM_002215	Homo sapiens inter-aipha (globulin) inmotor, 111 polypopulae (11211), 1121
NM_000889	Homo sapiens integrin, beta 7 (ITGB7), mRNA
NM_002212	Homo sapiens integrin beta 4 binding protein (ITGB4BP), mRNA
NM_000213	Homo sapiens integrin, beta 4 (ITGB4), mRNA
NM_002211	Homo sapiens integrin, beta 1 (fibronectin receptor, beta polypeptide, antigen CD29 includes MDF2, MSK12) (ITGB1), mRNA
NM_002210	Homo sapiens integrin, alpha V (vitronectin receptor, alpha polypeptide, antiger CD51) (ITGAV), mRNA
NM_002209	Homo sapiens integrin, alpha L (antigen CD11A (p180), lymphocyte function-associated antigen 1; alpha polypeptide) (ITGAL), mRNA
NM 002206	Homo sapiens integrin, alpha 7 (ITGA7), mRNA
NM_002205	Homo sapiens integrin, alpha 5 (fibronectin receptor, alpha polypeptide) (ITGA5), mRNA
NM 003749	Homo sapiens insulin receptor substrate 2 (IRS2), mRNA
NM 001571	Homo sapiens interferon regulatory factor 3 (IRF3), mRNA
NM 002198	Homo sapiens interferon regulatory factor 1 (IRF1), mRNA

	TY : i i i i i i i i i i i i i i i i i i
NM_002196	Homo sapiens insulinoma-associated 1 (INSM1), mRNA
NM_002195	Homo sapiens insulin-like 4 (placenta) (INSL4), mRNA
NM_001565	Homo sapiens small inducible cytokine subfamily B (Cys-X-Cys), member 10 (SCYB10), mRNA
NM_002192	Homo sapiens inhibin, beta A (activin A, activin AB alpha polypeptide)
	(INHBA), mRNA
NM_001564	Homo sapiens inhibitor of growth family, member 1-like (ING1L), mRNA
NM_003669	Homo sapiens inactivation escape 1 (INE1), mRNA
NM_000884	Homo sapiens IMP (inosine monophosphate) dehydrogenase 2 (IMPDH2), mRNA
NM_000883	Homo sapiens IMP (inosine monophosphate) dehydrogenase 1 (IMPDH1), mRNA
NM 001557	Homo sapiens interleukin 8 receptor, beta (IL8RB), mRNA
NM 000634	Homo sapiens interleukin 8 receptor, alpha (IL8RA), mRNA
NM 002185	Homo sapiens interleukin 7 receptor (IL7R), mRNA
NM 000880	Homo sapiens interleukin 7 (IL7), mRNA
NM 002184	Homo sapiens interleukin 6 signal transducer (gp130, oncostatin M receptor)
14141_002164	(IL6ST), mRNA
NM 000565	Homo sapiens interleukin 6 receptor (IL6R), mRNA
NM 000879	Homo sapiens interleukin 5 (colony-stimulating factor, eosinophil) (IL5), mRNA
NM 000589	Homo sapiens interleukin 4 (IL4), mRNA
NM 000588	Homo sapiens interleukin 3 (colony-stimulating factor, multiple) (IL3), mRNA
NM 000878	Homo sapiens interleukin 2 receptor, beta (IL2RB), mRNA
NM 003854	Homo sapiens interleukin 1 receptor-like 2 (IL1RL2), mRNA
NM 002182	Homo sapiens interleukin 1 receptor accessory protein (IL1RAP), mRNA
NM 000877	Homo sapiens interleukin 1 receptor, type I (IL1R1), mRNA
NM 003853	Homo sapiens interleukin 18 receptor accessory protein (IL18RAP), mRNA
NM 003855	Homo sapiens interleukin 18 receptor 1 (IL18R1), mRNA
NM 001562	Homo sapiens interleukin 18 (interferon-gamma-inducing factor) (IL18), mRNA
NM 002190	Homo sapiens interleukin 17 (cytotoxic T-lymphocyte-associated serine esterase
NWI_002190	8) (IL17), mRNA
NM_002189	Homo sapiens interleukin 15 receptor, alpha (IL15RA), mRNA
NM_002188	Homo sapiens interleukin 13 (IL13), mRNA
NM_001559	Homo sapiens interleukin 12 receptor, beta 2 (IL12RB2), mRNA
NM_002187	Homo sapiens interleukin 12B (natural killer cell stimulatory factor 2, cytotoxic lymphocyte maturation factor 2, p40) (IL12B), mRNA
ND ( 000000	Homo sapiens interleukin 12A (natural killer cell stimulatory factor 1, cytotoxic
NM_000882	lymphocyte maturation factor 1, p35) (IL12A), mRNA
ND 4 000629	Homo sapiens interleukin 10 receptor, beta (IL10RB), mRNA
NM_000628	Homo sapiens interleukin 10 receptor, alpha (IL10RA), mRNA
NM_001558 NM_003639	Homo sapiens inhibitor of kappa light polypeptide gene enhancer in B-cells,
	kinase gamma (IKBKG), mRNA
NM_003640	Homo sapiens inhibitor of kappa light polypeptide gene enhancer in B-cells, kinase complex-associated protein (IKBKAP), mRNA
NM 001542	Homo sapiens immunoglobulin superfamily, member 3 (IGSF3), mRNA
NM_001542 NM_001555	Homo sapiens immunoglobulin superfamily, member 1 (IGSF1), mRNA
	Homo sapiens immunoglobulin superfamily, memoer 1 (1001 1), mRVA  Homo sapiens immunoglobulin mu binding protein 2 (IGHMBP2), mRNA
NM_002180	Homo sapiens immunoglobulin mu binding protein 2 (1011WB1 2), mkdvA  Homo sapiens insulin-like growth factor binding protein 7 (IGFBP7), mRNA
NM_001553	Homo sapiens insulin-like growth factor binding protein 7 (IGFBF7), mRNA  Homo sapiens insulin-like growth factor binding protein 3 (IGFBP3), mRNA
NM_000598	Homo sapiens insulin-like growth factor binding protein 3 (IGFBF3), INCVA  Homo sapiens insulin-like growth factor binding protein 1 (IGFBP1), mRNA
NM_000596	Tomo sapiens insulin-like growth factor billiang protein 1 (IOPDI 1), filking
NM_001554	Homo sapiens cysteine-rich, angiogenic inducer, 61 (CYR61), mRNA  Homo sapiens insulin-like growth factor 2 receptor (IGF2R), mRNA
NM_000876_	nomo sapiens insumi-like grown factor 2 feceptor (1012k), findax

	1 (IDDD1) DNA
NM 001550	Homo sapiens interferon-related developmental regulator 1 (IFRD1), mRNA
000155	TI are remined interferon omega (IFNWI), IIINNA
2 1212	C 1 fibroblact (IENRI) MKNA
ND ( 000974	Homo saniens interferon (alpha, beta and omega) receptor 2 (HTML2), made a
ND4 002170	Homo saniens interferon, alpha 8 (IFNA8), mknA
NR ( 000160	Homo saniens interferon, alpha 5 (IFNA5), mRNA
NM 002175	Homo saniens interferon, alpha 21 (IFNAZI), mRNA
NM 002173	Homo saniens interferon, alpha 16 (IFNA16), mknA
NM 002172	Home saniens interferon, alpha 14 (IFNA14), mKNA
NM 002171	
NM_001549	Homo sapiens interferon-induced protein with terral copeptide repeats.
NM_001548	Homo sapiens interferon-induced protein with tetratricopeptide repeats 1 (H-111),
NM_003641	mRNA Homo sapiens interferon induced transmembrane protein 1 (9-27) (IFITM1),
	mRNA  I factor (complement) (IF) mRNA
NM_000204	Homo sapiens I factor (complement) (IF), mRNA  Homo sapiens isocitrate dehydrogenase 2 (NADP+), mitochondrial (IDH2),
NM_002168	1'
	nuclear gene encoding mitochondrial protein, mRNA  Homo sapiens inhibitor of DNA binding 4, dominant negative helix-loop-helix
NM_001546	
	protein (ID4), mRNA Homo sapiens inhibitor of DNA binding 2, dominant negative helix-loop-helix
NM_002166	
	protein (ID2), mRNA Homo sapiens inhibitor of DNA binding 1, dominant negative helix-loop-helix
NM_002165	. : (TD1) DNIA
	protein (ID1), mRNA  Homo sapiens hexabrachion (tenascin C, cytotactin) (HXB), mRNA
NM_002160	t f 1 1
NM_000871	· r 1 - 1 - 1 - 1 - 1 - 1 - 1 - 1 - 1 - 1
NM_000869	5 1-1-department on the (Serotonin) (Section 40 (111400))
NM_000868	
NM_000867	F landwaystermine (serotonini tecepiol 12 (1111112))
NM_000865	Transferred F budgovutning (Seroloni)) I Coopiul ID (111112)
NM_000864	F hydroxytromtomine (serotonin) (cuculot 1D (1111(1D))
NM_000863	Homo sapiens 5-hydroxytryptamine (serotonin) receptor 1A (HTR1A), mRNA  Homo sapiens 5-hydroxytryptamine (serotonin) receptor 1A (HTR1A), mRNA
NM_000524	1'-4-4'- 1 (HTN1) mPNA
NM_002159	Homo sapiens histatin 1 (HTN1), mixtyA  Homo sapiens human T-cell leukemia virus enhancer factor (HTLF), mRNA
NM_002158	Homo sapiens heat shock 27kD protein 2 (HSPB2), mRNA  Homo sapiens heat shock 27kD protein 2 (HSPB2), mRNA
NM_001541	Homo sapiens heat shock 70kD protein 6 (HSP70B') (HSPA6), mRNA  Homo sapiens heat shock 70kD protein 6 (HSP70B') (HSPA6), mRNA
NM_002155	Total and a material DNA Like 2 (HS12), INKNA
NM_001539	Homo sapiens heat snock protein, Division 2 (1983), Homo sapiens hydroxy-delta-5-steroid dehydrogenase, 3 beta- and steroid delta-
NM_000198	
NT 6 000063	isomerase 2 (HSD3B2), mRNA  Homo sapiens hydroxy-delta-5-steroid dehydrogenase, 3 beta- and steroid delta-
NM_000862	1 (TCD2D1) mDNA
ND 4 000414	hardrowystoroid (17-beta) dehydrogenase 4 (IISD17D4), illiaux
NM_000414	in a hydroxy storoid (17-heta) dehydrogenase 2 (113D1/D2), indiviz
NM_002153	- 1decreased (17 beta) denvertopellase i (11001/101); indus-
NM_000413	in a by drown of the fall denvertogenase 2 (110D 110D); machine
NM_000196	1 //head professe settile [1111111]
NM_002151	Tr
NM_000860	
NM_002150	
NM_002143	
NM_002148	Tionio saprens nomes con 210 (120-

NM_002147	Homo sapiens homeo box B5 (HOXB5), mRNA
NM_002146	Homo sapiens homeo box B3 (HOXB3), mRNA
NM_002145	Homo sapiens homeo box B2 (HOXB2), mRNA
NM_002144	Homo sapiens homeo box B1 (HOXB1), mRNA
NM_002142	Homo sapiens homeo box A9 (HOXA9), mRNA
NM_002141	Homo sapiens homeo box A4 (HOXA4), mRNA
NM_000522	Homo sapiens homeo box A13 (HOXA13), mRNA
NM_002139	Homo sapiens RNA binding motif protein, X chromosome (RBMX), mRNA
NM_000457	Homo sapiens hepatocyte nuclear factor 4, alpha (HNF4A), mRNA
NM_002135	Homo sapiens nuclear receptor subfamily 4, group A, member 1 (NR4A1), mRNA
NM 002133	Homo sapiens heme oxygenase (decycling) 1 (HMOX1), mRNA
NM_002131	Homo sapiens high-mobility group (nonhistone chromosomal) protein isoforms I and Y (HMGIY), mRNA
NM_002130	Homo sapiens 3-hydroxy-3-methylglutaryl-Coenzyme A synthase 1 (soluble) (HMGCS1), mRNA
NM_002128	Homo sapiens high-mobility group (nonhistone chromosomal) protein 1 (HMG1), mRNA
NM 000190	Homo sapiens hydroxymethylbilane synthase (HMBS), mRNA
NM 002126	Homo sapiens hepatic leukemia factor (HLF), mRNA
NM_001531	Homo sapiens major histocompatibility complex, class I-like sequence (HLALS), mRNA
NM 002127	Homo sapiens HLA-G histocompatibility antigen, class I, G (HLA-G), mRNA
NM_002123	Homo sapiens major histocompatibility complex, class II, DQ beta 1 (HLA-DQB1), mRNA
NM_001530	Homo sapiens hypoxia-inducible factor 1, alpha subunit (basic helix-loop-helix transcription factor) (HIF1A), mRNA
NM 001528	Homo sapiens HGF activator (HGFAC), mRNA
NM_000187	Homo sapiens homogentisate 1,2-dioxygenase (homogentisate oxidase) (HGD), mRNA
NM 000410	Homo sapiens hemochromatosis (HFE), mRNA
NM 000186	Homo sapiens H factor 1 (complement) (HF1), mRNA
NM 003865	Homo sapiens homeo box (expressed in ES cells) 1 (HESX1), mRNA
NM 002112	Homo sapiens histidine decarboxylase (HDC), mRNA
NM 002110	Homo sapiens hemopoietic cell kinase (HCK), mRNA
NM 003642	Homo sapiens histone acetyltransferase 1 (HAT1), mRNA
NM 001523	Homo sapiens hyaluronan synthase 1 (HAS1), mRNA
NM_000183	Homo sapiens hydroxyacyl-Coenzyme A dehydrogenase/3-ketoacyl-Coenzyme A thiolase/enoyl-Coenzyme A hydratase (trifunctional protein), beta subunit (HADHB), mRNA
NM_000182	Homo sapiens hydroxyacyl-Coenzyme A dehydrogenase/3-ketoacyl-Coenzyme A thiolase/enoyl-Coenzyme A hydratase (trifunctional protein), alpha subunit (HADHA), mRNA
NM_003548	Homo sapiens H4 histone, family 2 (H4F2), mRNA
NM_003547	Homo sapiens H4 histone family, member L (H4FL), mRNA
NM_003544	Homo sapiens H4 histone family, member I (H4FI), mRNA
NM_003493	Homo sapiens H3 histone family, member T (H3FT), mRNA
NM 003537	Homo sapiens H3 histone family, member L (H3FL), mRNA
NM 003534	Homo sapiens H3 histone family, member H (H3FH), mRNA
NM 003532	Homo sapiens H3 histone family, member D (H3FD), mRNA
NM 003531	Homo sapiens H3 histone family, member C (H3FC), mRNA
NM 003530	Homo sapiens H3 histone family, member B (H3FB), mRNA
	1 7,

NM 003529	Homo sapiens H3 histone family, member A (H3FA), mRNA
NIM 002107	Homo saniens H3 histone, family 3A (H3F3A), mRNA
NM 003528	Homo sapiens H2B histone family, member Q (H2BFQ), mRNA
NM 003526	Homo sapiens H2B histone family, member L (H2BFL), mRNA
NM 003525	Homo saniens H2B histone family, member K (H2BFK), mRNA
NM 003524	Homo sapiens H2B histone family, member J (H2BFJ), mRNA
NM 003523	Homo sapiens H2B histone family, member H (H2BFH), mRNA
NM 003522	Homo saniens H2B histone family, member G (H2BFG), mRNA
VM 003518	Home saniens H2B histone family, member A (H2BFA), mRNA
NM 002106	Homo saniens H2A histone family, member Z (H2AFZ), mRNA
NM 003516	Homo saniens H2A histone family, member O (H2AFO), mRNA
NM 003513	Homo saniens H2A histone family, member M (H2AFM), mRNA
NM 003512	Home ganiens H2A histone family member L (H2AFL), mRNA
NM 003612	Homo sapiens sema domain, immunoglobulin domain (1g), and GPI mentorate
1417_005012	emphor (semanhorin) 7A (SEMA7A), mRNA
NM_002104	Homo sapiens granzyme K (serine protease, granzyme 3; tryptase II) (GZMK),
1111_00210	mRNA
NM 002103	Homo sapiens glycogen synthase 1 (muscle) (GYS1), mRNA
NM 002102	Homo sapiens glycophorin E (GYPE), mRNA
NM 000181	Homo sapiens glucuronidase, beta (GUSB), mRNA
NM 000858	Homo saniens quanylate kinase 1 (GUK1), mRNA
NM 001522	Home conjens granylate cyclase 2F, retinal (GUCY2F), mRNA
NM_000180	Homo sapiens guanylate cyclase 2D, membrane (retina-specific) (GUCY2D),
14141_000100	mDNA
NM 000857	Homo saniens quanylate cyclase 1, soluble, beta 3 (GUCY1B3), mRNA
NM 000856	Homo conjens quanylate cyclase 1, soluble, alpha 3 (GUC I IAS), Ilicia
NM 000855	Home copiens quanylate cyclase 1 soluble, alpha 2 (GUCY1A2), mknA
NM 000409	Homo conjens quanylate cyclase activator IA (retina) (GUCAIA), IIIAVA
NM_001517	Homo sapiens general transcription factor IIH, polypeptide 4 (32kD subunit)
	(GTF2H4), mRNA
NM_002096	Homo sapiens general transcription factor IIF, polypeptide 1 (74kD subunit)
	(GTF2F1), mRNA
NM_002095	Homo sapiens general transcription factor IIE, polypeptide 2 (beta subunit,
	34kD) (GTF2E2), mRNA
NM_001513	Homo sapiens glutathione transferase zeta 1 (maleylacetoacetate isomerase)
	(GSTZ1), mRNA
NM_000853	Homo sapiens glutathione S-transferase theta 1 (GSTT1), mRNA
NM_000851	Homo sapiens glutathione S-transferase M5 (GSTM5), mRNA
NM_000850	Homo sapiens glutathione S-transferase M4 (GSTM4), mRNA  Homo sapiens glutathione S-transferase M3 (brain) (GSTM3) mRNA
NM_000849	Homo sapiens glutathione S-transferase M3 (brain) (GSTM3), mRNA  Homo sapiens glutathione S-transferase M3 (brain) (GSTM2), mRNA
NM_000848	Homo sapiens glutathione S-transferase M2 (muscle) (GSTM2), mRNA  Homo sapiens glutathione S-transferase M2 (muscle) (GSTM2), mRNA
NM_001512	Homo sapiens glutathione S-transferase A4 (GSTA4), mRNA
NM_000846	Homo sapiens glutathione S-transferase A2 (GSTA2), mRNA
NM_000178	Homo sapiens glutathione synthetase (GSS), mRNA
NM_002094	Homo sapiens G1 to S phase transition 1 (GSPT1), mRNA
NM_000177	Homo sapiens gelsolin (amyloidosis, Finnish type) (GSN), mRNA
NM_002093	Homo sapiens glycogen synthase kinase 3 beta (GSK3B), mRNA
NM_002092	Homo saniens G-rich RNA sequence binding factor I (GRSFI), HIKNA
NM_002091	Homo sapiens gastrin-releasing peptide (GRP), mRNA
NM_002090	Homo sapiens GRO3 oncogene (GRO3), mRNA
NA 000000	Homo sapiens GRO2 oncogene (GRO2), mRNA
NM_002089	Homo sapiens GRO1 oncogene (melanoma growth stimulating activity, alpha)

	(GRO1), mRNA
ND 4 002007	
NM_002087	Homo sapiens granulin (GRN), mRNA  Homo sapiens glutamate receptor, metabotropic 8 (GRM8), mRNA
NM_000845	Homo sapiens glutamate receptor, inclauditopic o (GRWo), inclauditopic o (GRWo), inclauditopic o (GRWo), mPNA
NM_000844	Homo sapiens glutamate receptor, metabotropic 7 (GRM7), mRNA
NM_000841	Homo sapiens glutamate receptor, metabotropic 4 (GRM4), mRNA
NM_000840	Homo sapiens glutamate receptor, metabotropic 3 (GRM3), mRNA
NM_000176	Homo sapiens nuclear receptor subfamily 3, group C, member 1 (NR3C1), mRNA
NM_000831	Homo sapiens glutamate receptor, ionotropic, kainate 3 (GRIK3), mRNA
NM_000830	Homo sapiens glutamate receptor, ionotropic, kainate 1 (GRIK1), mRNA
NM_002086	Homo sapiens growth factor receptor-bound protein 2 (GRB2), mRNA
NM_002085	Homo sapiens glutathione peroxidase 4 (phospholipid hydroperoxidase) (GPX4), mRNA
NM 002083	Homo sapiens glutathione peroxidase 2 (gastrointestinal) (GPX2), mRNA
NM 002082	Homo sapiens G protein-coupled receptor kinase 6 (GPRK6), mRNA
NM 001504	Homo sapiens G protein-coupled receptor 9 (GPR9), mRNA
NM 001508	Homo sapiens G protein-coupled receptor 39 (GPR39), mRNA
NM 001507	Homo sapiens G protein-coupled receptor 38 (GPR38), mRNA
NM 001506	Homo sapiens G protein-coupled receptor 32 (GPR32), mRNA
NM 001505	Homo sapiens G protein-coupled receptor 32 (GrR32), mRNA
NM 001503	Homo sapiens glycosylphosphatidylinositol specific phospholipase D1 (GPLD1),
	mRNA
NM_000408	Homo sapiens glycerol-3-phosphate dehydrogenase 2 (mitochondrial) (GPD2), mRNA
NM_001448	Homo sapiens glypican 4 (GPC4), mRNA
NM 002081	Homo sapiens glypican 1 (GPC1), mRNA
NM 000174	Homo sapiens glycoprotein IX (platelet) (GP9), mRNA
NM 000173	Homo sapiens glycoprotein Ib (platelet), alpha polypeptide (GP1BA), mRNA
NM_002080	Homo sapiens glutamic-oxaloacetic transaminase 2, mitochondrial (aspartate aminotransferase 2) (GOT2), nuclear gene encoding mitochondrial protein, mRNA
NM_002079	Homo sapiens glutamic-oxaloacetic transaminase 1, soluble (aspartate aminotransferase 1) (GOT1), mRNA
NM_002076	Homo sapiens glucosamine (N-acetyl)-6-sulfatase (Sanfilippo disease IIID) (GNS), mRNA
NM 001501	Homo sapiens gonadotropin-releasing hormone 2 (GNRH2), mRNA
NM 000825	Homo sapiens gonadotropin-releasing hormone 1 (leutinizing-releasing
	hormone) (GNRH1), mRNA
NM 002075	Homo sapiens guanine nucleotide binding protein (G protein), beta polypeptide 3
1111_002073	(GNB3), mRNA
NM 002073	Homo sapiens guanine nucleotide binding protein (G protein), alpha z
14141_002073	polypeptide (GNAZ), mRNA
NM 000172	Homo sapiens guanine nucleotide binding protein (G protein), alpha transducing
NWI_000172	activity polypeptide 1 (GNAT1), mRNA
ND ( 002072	
NM_002072	Homo sapiens guanine nucleotide binding protein (G protein), q polypeptide (GNAQ), mRNA
NM_002071	Homo sapiens guanine nucleotide binding protein (G protein), alpha activating activity polypeptide, olfactory type (GNAL), mRNA
NM_002070	Homo sapiens guanine nucleotide binding protein (G protein), alpha inhibiting
_	activity polypeptide 2 (GNAI2), mRNA
NM 002068	Homo sapiens guanine nucleotide binding protein (G protein), alpha 15 (Gq
	class) (GNA15), mRNA
L	

	- 1 11 1 in time protein (G protein) alpha 11 (Go
NM_002067	Homo sapiens guanine nucleotide binding protein (G protein), alpha 11 (Gq
	class) (GNA11), mRNA
NM_003875	Homo sapiens guanine monphosphate synthetase (GMPS), mRNA
NM_002066	Homo sapiens GPI anchored molecule like protein (GML), mRNA
NM_001500	Homo sapiens GDP-mannose 4,6-dehydratase (GMDS), mRNA
NM_002065	Homo sapiens glutamate-ammonia ligase (glutamine synthase) (GLUL), mRNA
NM_002064	Homo sapiens glutaredoxin (thioltransferase) (GLRX), mRNA
NM_000824	Homo sapiens glycine receptor, beta (GLRB), mRNA
NM_002063	Homo sapiens glycine receptor, alpha 2 (GLRA2), mRNA
NM_002062	Homo sapiens glucagon-like peptide 1 receptor (GLP1R), mRNA
NM_000170	Homo sapiens glycine dehydrogenase (decarboxylating; glycine decarboxylase,
	glycine cleavage system protein P) (GLDC), mRNA
NM_000169	Homo sapiens galactosidase, alpha (GLA), mRNA
NM_000167	Homo sapiens glycerol kinase (GK), mRNA
NM_000166	Homo sapiens gap junction protein, beta 1, 32kD (connexin 32, Charcot-Marie-
_	Tooth neuropathy, X-linked) (GJB1), mRNA
NM 002060	Homo sapiens gap junction protein, alpha 4, 37kD (connexin 37) (GJA4), mRNA
NM_000164	Homo sapiens gastric inhibitory polypeptide receptor (GIPR), mRNA
NM 000823	Homo sapiens growth hormone releasing hormone receptor (GHRHR), mRNA
NM 000163	Homo sapiens growth hormone receptor (GHR), mRNA
NM 000821	Homo sapiens gamma-glutamyl carboxylase (GGCX), mRNA
NM 001495	Homo sapiens GDNF family receptor alpha 2 (GFRA2), mRNA
NM 002055	Homo sapiens glial fibrillary acidic protein (GFAP), mRNA
NM 003943	Homo sapiens genethonin 1 (GENX-3414), mRNA
NM 000514	Homo sapiens glial cell derived neurotrophic factor (GDNF), mRNA
NM 001493	Homo sapiens GDP dissociation inhibitor 1 (GDI1), mRNA
NM_001491	Homo sapiens glucosaminyl (N-acetyl) transferase 2, I-branching enzyme
	(GCNT2), mRNA
NM_001490	Homo sapiens glucosaminyl (N-acetyl) transferase 1, core 2 (beta-1,6-N-
	acetylglucosaminyltransferase) (GCNT1), mRNA
NM_000160	Homo sapiens glucagon receptor (GCGR), mRNA
NM_002054	Homo sapiens glucagon (GCG), mRNA
NM_001485	Homo sapiens gastrulation brain homeo box 2 (GBX2), mRNA
NM_001483	Homo sapiens glioblastoma amplified sequence (GBAS), mRNA
NM_002048	Homo sapiens growth arrest-specific 1 (GAS1), mRNA
NM 001481	Homo sapiens growth arrest-specific 11 (GAS11), mRNA
NM_000819	Homo sapiens phosphoribosylglycinamide formyltransferase,
	phosphoribosylglycinamide synthetase, phosphoribosylaminoimidazole
	synthetase (GART), mRNA
NM 002045	Homo sapiens growth associated protein 43 (GAP43), mRNA
NM 003614	Homo sapiens galanin receptor 3 (GALR3), mRNA
NM 000154	Homo sapiens galactokinase 1 (GALK1), mRNA
NM 001477	Homo sapiens G antigen 7B (GAGE7B), mRNA
NM 001476	Homo sapiens G antigen 6 (GAGE6), mRNA
NM 001475	Homo sapiens G antigen 5 (GAGE5), mRNA
NM 001474	Homo sapiens G antigen 4 (GAGE4), mRNA
NM 001473	Homo sapiens G antigen 3 (GAGE3), mRNA
NM 001472	Homo sapiens G antigen 2 (GAGE2), mRNA
NM 001468	Homo sapiens G antigen 1 (GAGE1), mRNA
NM 000818	Homo sapiens glutamate decarboxylase 2 (pancreatic islets and brain, 65kD)
	(GAD2), mRNA
NM 002043	Homo sapiens gamma-aminobutyric acid (GABA) receptor, rho 2 (GABRR2),

	mRNA
NM_002042	Homo sapiens gamma-aminobutyric acid (GABA) receptor, rho 1 (GABRR1),
NW1_002042	mRNA
NM_000402	Homo sapiens glucose-6-phosphate dehydrogenase (G6PD), nuclear gene
1111_000402	encoding mitochondrial protein, mRNA
NM 001469	Homo sapiens thyroid autoantigen 70kD (Ku antigen) (G22P1), mRNA
NM 002037	Homo sapiens FYN oncogene related to SRC, FGR, YES (FYN), mRNA
NM 002036	Homo sapiens Duffy blood group (FY), mRNA
NM 002035	Homo sapiens follicular lymphoma variant translocation 1 (FVT1), mRNA
NM_000150	Homo sapiens fucosyltransferase 6 (alpha (1,3) fucosyltransferase) (FUT6),
NM_000130	mRNA
NM_002034	Homo sapiens fucosyltransferase 5 (alpha (1,3) fucosyltransferase) (FUT5),
1111_002054	mRNA
NM_002033	Homo sapiens fucosyltransferase 4 (alpha (1,3) fucosyltransferase, myeloid-
11111_002033	specific) (FUT4), mRNA
NM_000149	Homo sapiens fucosyltransferase 3 (galactoside 3(4)-L-fucosyltransferase, Lewis
14141_000145	blood group included) (FUT3), mRNA
NM 000511	Homo sapiens fucosyltransferase 2 (secretor status included) (FUT2), mRNA
NM 000311	Homo sapiens fucosyltransferase 1 (galactoside 2-alpha-L-fucosyltransferase,
14141_000140	Bombay phenotype included) (FUT1), mRNA
NM 000147	Homo sapiens fucosidase, alpha-L- 1, tissue (FUCA1), mRNA
NM 002032	Homo sapiens ferritin, heavy polypeptide 1 (FTH1), mRNA
NM 000145	Homo sapiens follicle stimulating hormone receptor (FSHR), mRNA
NM 000510	Homo sapiens follicle stimulating hormone, beta polypeptide (FSHB), mRNA
NM 001463	Homo sapiens frizzled-related protein (FRZB), mRNA
NM 000144	Homo sapiens Friedreich ataxia (FRDA), mRNA
NM 001462	Homo sapiens formyl peptide receptor-like 1 (FPRL1), mRNA
NM 002029	Homo sapiens formyl peptide receptor 1 (FPR1), mRNA
NM 003838	Homo sapiens fucose-1-phosphate guanylyltransferase (FPGT), mRNA
NM 002027	Homo sapiens farnesyltransferase, CAAX box, alpha (FNTA), mRNA
NM 002025	Homo sapiens fragile X mental retardation 2 (FMR2), mRNA
NM 002024	Homo sapiens fragile X mental retardation 1 (FMR1), mRNA
NM 001461	Homo sapiens flavin containing monooxygenase 5 (FMO5), mRNA
NM 002022	Homo sapiens flavin containing monooxygenase 4 (FMO4), mRNA
	Homo sapiens flavin containing monooxygenase 2 (FMO2), mRNA
NM_001460 NM_002021	Homo sapiens flavin containing monooxygenase 2 (FMO2), mRNA  Homo sapiens flavin containing monooxygenase 1 (FMO1), mRNA
	Homo sapiens fms-related tyrosine kinase 4 (FLT4), mRNA
NM_002020	Homo sapiens fms-related tyrosine kinase 3 ligand (FLT3LG), mRNA
NM_001459	Homo sapiens fms-related tyrosine kinase 3 figand (TET3EO), find the Homo sapiens fms-related tyrosine kinase 1 (vascular endothelial growth
NM_002019	factor/vascular permeability factor receptor) (FLT1), mRNA
NIM 001455	Homo sapiens forkhead box O3A (FOXO3A), mRNA
NM_001455	Homo sapiens forkhead box O3A (FOXO3A), mRNA  Homo sapiens forkhead box C1 (FOXC1), mRNA
NM_001453	Homo sapiens forkhead box C1 (FOXC1), mRNA  Homo sapiens forkhead box F1 (FOXF1), mRNA
NM_001451	
NM_001450	Homo sapiens four and a half LIM domains 2 (FHL2), mRNA Homo sapiens four and a half LIM domains 1 (FHL1), mRNA
NM_001449	
NM_002012	Homo sapiens fragile histidine triad gene (FHIT), mRNA
NM_000143	Homo sapiens fumarate hydratase (FH), mRNA
NM_002002	Homo sapiens Fc fragment of IgE, low affinity II, receptor for (CD23A) (FCER2), mRNA
NM 002001	Homo sapiens Fc fragment of IgE, high affinity I, receptor for; alpha polypeptide
	(FCER1A), mRNA
NM_002000	Homo sapiens Fc fragment of IgA, receptor for (FCAR), mRNA

Iomo sapiens fructose-1,6-bisphosphatase 2 (FBP2), mRNA Iomo sapiens fibulin 2 (FBLN2), mRNA Iomo sapiens forkhead box H1 (FOXH1), mRNA
Is no serious forkhead how H1 (FOXH1), mRNA
Iomo sapiens forkhead box H1 (FOXH1), mRNA
Iomo sapiens coagulation factor II (thrombin) receptor-like 3 (F2RL3), mRNA
Iomo sapiens SH2 domain protein 2A (SH2D2A), mRNA
Iomo sapiens exostoses (multiple)-like 3 (EXTL3), mRNA
Jomo conjens envonlakin (EVPL) mKNA
Iomo seniens electron-transfer-flavoprotein, beta polypeptide (EIFB), IIIKINA
I) (FTFA) nuclear gene encoding mitochondrial protein, mRNA
Homo sapiens estrogen-related receptor gamma (ESRRG), mRNA
Jome conjens estrogen recentor 1 (ESR1), mRNA
Temps comions evoision repair cross-complementing rodent repair deficiency,
complementation group 5 (xeroderma pigmentosum, complementation group G (Cockayne syndrome)) (ERCC5), mRNA
Homo sapiens excision repair cross-complementing rodent repair deficiency,
complementation group 1 (includes overlapping antisense sequence) (ERCC1), mRNA
Home conjune equipophil peroxidase (EPX), mRNA
Homo sapiens epidermal growth factor receptor pathway substrate 15 (EPS15), mRNA
Homo sapiens erythropoietin (EPO), mRNA
Homo sapiens epimorphin (EPIM), mRNA
Homo sapiens erythrocyte membrane protein band 4.1-like 2 (EPB41L2), mRNA
Homo sapiens endothelial PAS domain protein 1 (EPAS1), mRNA
Homo sapiens glutamyl aminopeptidase (aminopeptidase A) (ENPEP), mRNA
Homo sapiens grutamy animopeptidase (animopeptidase 15) (15) Homo sapiens egf-like module containing, mucin-like, hormone receptor-like
sequence 1 (EMR1), mRNA
Homo sapiens epithelial membrane protein 3 (EMP3), mRNA
Homo sapiens epithelial membrane protein 2 (EMP2), mRNA
Homo sapiens epithelial membrane protein 1 (EMP1), mRNA
Homo saniens E74-like factor 4 (ets domain transcription factor) (ELF4), IIIKNA
Homo sapiens ELAV (embryonic lethal, abnormal vision, Drosophila)-like I (Hu antigen R) (ELAVL1), mRNA
Homo saniens elastase 2. neutrophil (ELA2), mRNA
Homo saniens eukaryotic translation initiation factor 5A (EIF5A), mRNA
Homo sapiens eukaryotic translation initiation factor 4 gamma, 2 (EIF4G2),
Homo sapiens eukaryotic translation initiation factor 4E binding protein 3 (EIF4EBP3), mRNA
Homo sapiens eukaryotic translation initiation factor 4E (EIF4E), mRNA
Homo sapiens eukaryotic translation initiation factor 4A, isoform 1 (EIF4A1), mRNA
Homo sapiens eukaryotic translation initiation factor 3, subunit 7 (zeta, 66/67kD)
Homo sapiens eukaryotic translation initiation factor 3, subunit 6 (48kD)
Homo sapiens eukaryotic translation initiation factor 3, subunit 5 (epsilon, 4/kD)
Homo sapiens eukaryotic translation initiation factor 3, subunit 2 (beta, 36kD) (EIF3S2) mRNA
Homo sapiens eukaryotic translation initiation factor 3, subunit 10 (theta,

	150/170kD) (EIF3S10), mRNA
NM_001415	Homo sapiens eukaryotic translation initiation factor 2, subunit 3 (gamma, 52kD) (EIF2S3), mRNA
NM_003908	Homo sapiens eukaryotic translation initiation factor 2, subunit 2 (beta, 38kD) (EIF2S2), mRNA
NM_001966	Homo sapiens enoyl-Coenzyme A, hydratase/3-hydroxyacyl Coenzyme A dehydrogenase (EHHADH), nuclear gene encoding mitochondrial protein, mRNA
NM 001965	Homo sapiens early growth response 4 (EGR4), mRNA
NM 001964	Homo sapiens early growth response 1 (EGR1), mRNA
NM 001406	Homo sapiens ephrin-B3 (EFNB3), mRNA
NM 001962	Homo sapiens ephrin-A5 (EFNA5), mRNA
NM 001405	Homo sapiens ephrin-A2 (EFNA2), mRNA
NM 001961	Homo sapiens eukaryotic translation elongation factor 2 (EEF2), mRNA
NM_001958	Homo sapiens eukaryotic translation elongation factor 1 alpha 2 (EEF1A2), mRNA
NM 001956	Homo sapiens endothelin 2 (EDN2), mRNA
NM 001955	Homo sapiens endothelin 1 (EDN1), mRNA
NM_003775	Homo sapiens endothelial differentiation, G-protein-coupled receptor 6 (EDG6), mRNA
NM 001399	Homo sapiens ectodermal dysplasia 1, anhidrotic (ED1), mRNA
NM 001397	Homo sapiens endothelin converting enzyme 1 (ECE1), mRNA
NM 003240	Homo sapiens endometrial bleeding associated factor (left-right determination,
1414_005210	factor A; transforming growth factor beta superfamily) (EBAF), mRNA
NM 001948	Homo sapiens dUTP pyrophosphatase (DUT), mRNA
NM_001945	Homo sapiens diphtheria toxin receptor (heparin-binding epidermal growth factor-like growth factor) (DTR), mRNA
NM 001939	Homo sapiens dystrophin related protein 2 (DRP2), mRNA
NM_001938	Homo sapiens down-regulator of transcription 1, TBP-binding (negative cofactor 2) (DR1), mRNA
NM 001387	Homo sapiens dihydropyrimidinase-like 3 (DPYSL3), mRNA
NM 001385	Homo sapiens dihydropyrimidinase (DPYS), mRNA
NM_001935	Homo sapiens dipeptidylpeptidase IV (CD26, adenosine deaminase complexing protein 2) (DPP4), mRNA
NM_003863	Homo sapiens dolichyl-phosphate mannosyltransferase polypeptide 2, regulatory subunit (DPM2), mRNA
NM 001380	Homo sapiens dedicator of cyto-kinesis 1 (DOCK1), mRNA
NM 001379	Homo sapiens DNA (cytosine-5-)-methyltransferase 1 (DNMT1), mRNA
NM 001379	Homo sapiens deoxyribonuclease II, lysosomal (DNASE2), mRNA
NM 001374	Homo sapiens deoxyribonuclease I-like 2 (DNASE1L2), mRNA
NM 001934	Homo sapiens distal-less homeobox 4 (DLX4), mRNA
NM 001934	Homo sapiens dihydrolipoamide S-succinyltransferase (E2 component of 2-oxo-
14141_001955	glutarate complex) (DLST), mRNA
NM 001362	Homo sapiens deiodinase, iodothyronine, type III (DIO3), mRNA
NM 001360	Homo sapiens 7-dehydrocholesterol reductase (DHCR7), mRNA
NM_003670	Homo sapiens basic helix-loop-helix domain containing, class B, 2 (BHLHB2), mRNA
NM_001354	Homo sapiens aldo-keto reductase family 1, member C2 (dihydrodiol dehydrogenase 2; bile acid binding protein; 3-alpha hydroxysteroid dehydrogenase, type III) (AKR1C2), mRNA
NM_000790	Homo sapiens dopa decarboxylase (aromatic L-amino acid decarboxylase) (DDC), mRNA

	(1) 1 (ancietongin I converting enzyme)
NM_000789	Homo sapiens dipeptidyl carboxypeptidase 1 (angiotensin I converting enzyme)
	(ACE), mRNA
NM_001920	Homo sapiens decorin (DCN), mRNA
NM_000788	Homo sapiens deoxycytidine kinase (DCK), mRNA
NM_001919	Homo sapiens dedecenoyl-Coenzyme A delta isomerase (3,2 trans-enoyl-
	Coenzyme A isomerase) (DCI), mRNA  Homo sapiens dihydrolipoamide branched chain transacylase (E2 component of
NM_001918	Homo sapiens dihydrolipoamide branched chain transdoylase (22 composition branched chain keto acid dehydrogenase complex; maple syrup urine disease)
	branched chain keto acid denydrogenase complex, mapio syrap
	(DBT), mRNA Homo sapiens D site of albumin promoter (albumin D-box) binding protein
NM_001352	Homo sapiens D site of aroundin productor (arounding b sorry streams p
	(DBP), mRNA  Homo sapiens deleted in azoospermia-like (DAZL), mRNA
NM_001351	Homo sapiens death-associated protein 6 (DAXX), mRNA
NM_001350	Homo sapiens defender against cell death 1 (DAD1), mRNA
NM_001344	Homo sapiens DEK oncogene (DNA binding) (DEK), mRNA
NM_003472	Homo sapiens DEK oncogene (DIVY omething) (222), Homo sapiens cytochrome P450, subfamily IIIA (niphedipine oxidase),
NM_000776	Homo sapiens cytochrome 1450, subtaining that (Approved
	polypeptide 3 (CYP3A3), mRNA  Homo sapiens cytochrome c-1 (CYC1), mRNA
NM_001916	Homo sapiens cytochrome b-5 (CYB5), nuclear gene encoding mitochondrial
NM_001914	Homo sapiens cytochrome 0-3 (C 1 B3), national gono one of the control of the con
1 000000	Protein, mRNA Homo sapiens CAAX box 1 (CXX1), mRNA
NM_003928	Homo sapiens CAAA box 1 (CAST), media frame 5 (CXORF5), mRNA  Homo sapiens chromosome X open reading frame 5 (CXORF5), mRNA
NM_003611	Homo sapiens chemokine (C-X-C motif), receptor 4 (fusin) (CXCR4), mRNA
NM_003467	Homo sapiens coxsackie virus and adenovirus receptor (CXADR), mRNA
NM_001338	Homo sapiens coxsackie viius and addition viius receptor
NM_003478	Homo sapiens cullin 5 (CUL5), mRNA
NM_003591	Homo sapiens cullin 2 (CUL2), mRNA
NM_001336	Homo sapiens cathepsin Z (CTSZ), mRNA Homo sapiens cathepsin W (lymphopain) (CTSW), mRNA
NM_001335	Homo sapiens cattlepsin w (tymphopam) (CTSL), mRNA
NM_001912	Homo sapiens cathepsin L (CTSL), mRNA  Homo sapiens cathepsin L2 (CTSL2), mRNA
NM_001333	Homo sapiens cathepsin L2 (CTSL2), interview Homo sapiens cathepsin K (pycnodysostosis) (CTSK), mRNA
NM_000396	Homo sapiens cathepsin K (pychodysostosis) (CTSG), mRNA
NM_001911	Homo sapiens cathepsin G (CTSG), mRNA
NM_001910	Homo sapiens cathepsin E (CTSE), mRNA Homo sapiens cathepsin D (lysosomal aspartyl protease) (CTSD), mRNA
NM_001909	Homo sapiens cathersin C (CTSC) mRNA
NM_001814	Homo sapiens cathepsin C (CTSC), mRNA
NM_001908	Homo sapiens cathepsin B (CTSB), mRNA Homo sapiens chymotrypsin-like (CTRL), mRNA
NM_001907	Homo sapiens chymotrypsinogen B1 (CTRB1), mRNA
NM_001906	Homo sapiens CTP synthase (CTPS), mRNA
NM_001905	Homo sapiens CTF synthase (CTF 5), interest Homo sapiens catenin (cadherin-associated protein), beta 1 (88kD) (CTNNB1),
NM_001904	DATA
27.6.00500	mRNA Homo sapiens catenin (cadherin-associated protein), alpha-like 1 (CTNNAL1),
NM_003798	Homo sapiens catenin (cadierin-associated protein), aspira
77.5 001002	mRNA Homo sapiens catenin (cadherin-associated protein), alpha 1 (102kD)
NM_001903	(CTNNA1), mRNA
27.6 001000	Homo sapiens cystathionase (cystathionine gamma-lyase) (CTH), mRNA
NM_001902	Homo sapiens connective tissue growth factor (CTGF), mRNA
NM_001901	Homo sapiens cardiotrophin 1 (CTF1), mRNA
NM_001330	Homo sapiens cystatin B (stefin B) (CSTB), mRNA
NM_000100	Homo sapiens cystatin B (sterin B) (CSTB), internal Homo sapiens cystatin F (leukocystatin) (CST7), mRNA
NM_003650	Homo sapiens cystatin F (letikocystatin) (CST4), index- Homo sapiens cystatin E/M (CST6), mRNA
NM_001323	Homo sapiens cystatin D (CCT5), mRNA
NM_001900	Homo sapiens cystatin D (CST5), mRNA

NM 001899	Homo sapiens cystatin S (CST4), mRNA
NM_000099	Homo sapiens cystatin C (amyloid angiopathy and cerebral hemorrhage) (CST3),
14141_000033	mRNA
NM 001322	Homo sapiens cystatin SA (CST2), mRNA
NM 001898	Homo sapiens cystatin SN (CST1), mRNA
NM 001321	Homo sapiens cysteine and glycine-rich protein 2 (CSRP2), mRNA
NM 001896	Homo sapiens casein kinase 2, alpha prime polypeptide (CSNK2A2), mRNA
NM 001895	Homo sapiens casein kinase 2, alpha 1 polypeptide (CSNK2A1), mRNA
NM 001894	Homo sapiens casein kinase 1, epsilon (CSNK1E), mRNA
NM 001893	Homo sapiens casein kinase 1, delta (CSNK1D), mRNA
NM 001893	Homo sapiens casein kinase 1, alpha 1 (CSNK1A1), mRNA
NM 001892	Homo sapiens casein, beta (CSN2), mRNA
	Homo sapiens casein, beta (CSN2), mRNA  Homo sapiens casein, alpha (CSN1), mRNA
NM_001890	Homo sapiens colony stimulating factor 3 receptor (granulocyte) (CSF3R),
NM_000760	mRNA
ND4 000750	Homo sapiens colony stimulating factor 3 (granulocyte) (CSF3), mRNA
NM_000759	Homo sapiens colony stimulating factor 2 (granulocyte-macrophage) (CSF2),
NM_000758	mRNA
NIM 000757	Homo sapiens colony stimulating factor 1 (macrophage) (CSF1), mRNA
NM_000757	Homo sapiens cold shock domain protein A (CSDA), mRNA
NM_003651	Homo sapiens mitogen-activated protein kinase 14 (MAPK14), mRNA
NM_001315	Homo sapiens cartilage linking protein 1 (CRTL1), mRNA
NM_001884	Homo sapiens cartinage linking protein 1 (CRTE1), linking Homo sapiens collapsin response mediator protein 1 (CRMP1), mRNA
NM_001313	Homo sapiens conapsin response mediator protein 1 (CRVII 1), midV1
NM_001312	Homo sapiens cysteine-rich protein 2 (CRIP2), mRNA
NM_001311	Homo sapiens cysteine-rich protein 1 (intestinal) (CRIP1), mRNA
NM_000756	Homo sapiens corticotropin releasing hormone (CRH), mRNA
NM_001881	Homo sapiens cAMP responsive element modulator (CREM), mRNA
NM_003851	Homo sapiens cellular repressor of E1A-stimulated genes (CREG), mRNA
NM_001310	Homo sapiens cAMP responsive element binding protein-like 2 (CREBL2), mRNA
NM 001880	Homo sapiens activating transcription factor 2 (ATF2), mRNA
NM_003805	Homo sapiens CASP2 and RIPK1 domain containing adaptor with death domain
	(CRADD), mRNA
NM_001877	Homo sapiens complement component (3d/Epstein Barr virus) receptor 2 (CR2), mRNA
NM 000098	Homo sapiens carnitine palmitoyltransferase II (CPT2), nuclear gene encoding
1.2.2_00000	mitochondrial protein, mRNA
NM 001876	Homo sapiens carnitine palmitoyltransferase I, liver (CPT1A), nuclear gene
1111_001070	encoding mitochondrial protein, mRNA
NM 001875	Homo sapiens carbamoyl-phosphate synthetase 1, mitochondrial (CPS1), nuclear
1	gene encoding mitochondrial protein, mRNA
NM_000097	Homo sapiens coproporphyrinogen oxidase (coproporphyria, harderoporphyria)
112.2_00007	(CPO), mRNA
NM 001871	Homo sapiens carboxypeptidase B1 (tissue) (CPB1), mRNA
NM 001870	Homo sapiens carboxypeptidase A3 (mast cell) (CPA3), mRNA
NM 001869	Homo sapiens carboxypeptidase A2 (pancreatic) (CPA2), mRNA
NM 001868	Homo sapiens carboxypeptidase A1 (pancreatic) (CPA1), mRNA
NM 003571	Homo sapiens beaded filament structural protein 2, phakinin (BFSP2), mRNA
NM 001302	Homo sapiens cortistatin (CORT), mRNA
NM 003832	Homo sapiens contistatif (CORT), filed it  Homo sapiens phosphoserine phosphatase-like (PSPHL), mRNA
NM 001843	Homo sapiens contactin 1 (CNTN1), mRNA
NM_001843	Homo sapiens contactiff (CNTINT), mid.
19191 001042	110110 Sapiens emary neuronopino factor receptor (C11111), metri

	1 : 2 :11- (CNINI2) PNIA
	Homo sapiens calponin 3, acidic (CNN3), mRNA
NM_001299	Homo sapiens calponin 1, basic, smooth muscle (CNN1), mRNA
NM_001297	Homo sapiens cyclic nucleotide gated channel beta 1 (CNGB1), mRNA
NM_001298	Homo sapiens cyclic nucleotide gated channel alpha 3 (CNGA3), mRNA
NM_000087	Homo sapiens cyclic nucleotide gated channel alpha 1 (CNGA1), mRNA
NM_003570	Homo sapiens cytidine monophosphate-N-acetylneuraminic acid hydroxylase
	(CMP-N-acetylneuraminate monooxygenase) (CMAH), mRNA
NM_001836	Homo sapiens chymase 1, mast cell (CMA1), mRNA
NM_001831	Homo sapiens clusterin (complement lysis inhibitor, SP-40,40, sulfated
	glycoprotein 2, testosterone-repressed prostate message 2, apolipoprotein J)
	(CLU), mRNA
NM_001294	Homo sapiens cleft lip and palate associated transmembrane protein 1
	(CLPTM1), mRNA
NM_003476	Homo sapiens cysteine and glycine-rich protein 3 (cardiac LIM protein)
	(CSRP3), mRNA
NM_001293	Homo sapiens chloride channel, nucleotide-sensitive, 1A (CLNS1A), mRNA
NM_003277	Homo sapiens claudin 5 (transmembrane protein deleted in velocardiofacial
	syndrome) (CLDN5), mRNA
NM_001306	Homo sapiens claudin 3 (CLDN3), mRNA
NM_001829	Homo sapiens chloride channel 3 (CLCN3), mRNA
NM_001284	Homo sapiens adaptor-related protein complex 3, sigma 1 subunit (AP3S1),
	mRNA DATE OF COMMON PAIA
NM_001827	Homo sapiens CDC28 protein kinase 2 (CKS2), mRNA
NM_001826	Homo sapiens CDC28 protein kinase 1 (CKS1), mRNA
NM_001824	Homo sapiens creatine kinase, muscle (CKM), mRNA
NM_001823	Homo sapiens creatine kinase, brain (CKB), mRNA
NM_001281	Homo sapiens cytoskeleton-associated protein 1 (CKAP1), mRNA
NM_003613	Homo sapiens cartilage intermediate layer protein, nucleotide
	pyrophosphohydrolase (CILP), mRNA
NM_001278	Homo sapiens conserved helix-loop-helix ubiquitous kinase (CHUK), mRNA
NM_003654	Homo sapiens carbohydrate (chondroitin 6/keratan) sulfotransferase 1 (CHST1),
	mRNA (CVID) TO (
NM_000750	Homo sapiens cholinergic receptor, nicotinic, beta polypeptide 4 (CHRNB4),
	mRNA (CVIDADE)
NM_000749	Homo sapiens cholinergic receptor, nicotinic, beta polypeptide 3 (CHRNB3),
	mRNA
NM_000748	Homo sapiens cholinergic receptor, nicotinic, beta polypeptide 2 (neuronal)
	(CHRNB2), mRNA
NM_000746	Homo sapiens cholinergic receptor, nicotinic, alpha polypeptide 7 (CHRNA7),
	mRNA (CLIPNAS)
NM_000745	Homo sapiens cholinergic receptor, nicotinic, alpha polypeptide 5 (CHRNA5),
	mRNA (CYPRALA 4)
NM_000744	Homo sapiens cholinergic receptor, nicotinic, alpha polypeptide 4 (CHRNA4),
	mRNA
NM_000743	Homo sapiens cholinergic receptor, nicotinic, alpha polypeptide 3 (CHRNA3),
	mRNA
NM_000742	Homo sapiens cholinergic receptor, nicotinic, alpha polypeptide 2 (neuronal)
_	(CHRNA2), mRNA
NM_000741	Homo sapiens cholinergic receptor, muscarinic 4 (CHRM4), mRNA
NM_000740	Homo sapiens cholinergic receptor, muscarinic 3 (CHRM3), mRNA
NM 000739	Homo sapiens cholinergic receptor, muscarinic 2 (CHRM2), mRNA
NM 000738	Homo sapiens cholinergic receptor, muscarinic 1 (CHRM1), mRNA

	TI (CUNI) mPNA
NM_001822	Homo sapiens chimerin (chimaerin) 1 (CHN1), mRNA
NM_001821	Homo sapiens choroideremia-like (Rab escort protein 2) (CHML), mRNA
NM_001819	Homo sapiens chromogranin B (secretogranin 1) (CHGB), mRNA
NM_001269	Homo sapiens chromosome condensation 1 (CHC1), mRNA
NM_001267	Homo sapiens chondroadherin (CHAD), mRNA
NM_001817	Homo sapiens carcinoembryonic antigen-related cell adhesion molecule 4 (CEACAM4), mRNA
ND ( 001016	Homo sapiens carcinoembryonic antigen-related cell adhesion molecule 8
NM_001816	(CEACAM8), mRNA
NM_001815	Homo sapiens carcinoembryonic antigen-related cell adhesion molecule 3 (CEACAM3), mRNA
NM_003663	Homo sapiens CGG triplet repeat binding protein 1 (CGGBP1), mRNA
NM 001813	Homo sapiens centromere protein E (312kD) (CENPE), mRNA
NM_001808	Homo sapiens carboxyl ester lipase-like (bile salt-stimulated lipase-like) (CELL),
NM_001808	mRNA
NM_001807	Homo sapiens carboxyl ester lipase (bile salt-stimulated lipase) (CEL), mRNA
NM_001805	Homo sapiens CCAAT/enhancer binding protein (C/EBP), epsilon (CEBPE), mRNA
NM 001265	Homo sapiens caudal type homeo box transcription factor 2 (CDX2), mRNA
NM 001804	Homo sapiens caudal type homeo box transcription factor 1 (CDX1), mRNA
NM 001803	Homo sapiens CDW52 antigen (CAMPATH-1 antigen) (CDW52), mRNA
NM 001264	Homo sapiens corneodesmosin (CDSN), mRNA
NM 001263	Homo sapiens CDP-diacylglycerol synthase (phosphatidate cytidylyltransferase)
NWI_001203	1 (CDS1), mRNA
NM_001801	Homo sapiens cysteine dioxygenase, type I (CDO1), mRNA
NM_001769	Homo sapiens CD9 antigen (p24) (CD9), mRNA
NM_001768	Homo sapiens CD8 antigen, alpha polypeptide (p32) (CD8A), mRNA
NM_003874	Homo sapiens CD84 antigen (leukocyte antigen) (CD84), mRNA
NM_001781	Homo sapiens CD69 antigen (p60, early T-cell activation antigen) (CD69), mRNA
NM 001780	Homo sapiens CD63 antigen (melanoma 1 antigen) (CD63), mRNA
NM_001779	Homo sapiens CD58 antigen, (lymphocyte function-associated antigen 3)
ND 6 001770	(CD58), mRNA Homo sapiens CD48 antigen (B-cell membrane protein) (CD48), mRNA
NM_001778 NM_001777	Homo sapiens CD48 antigen (B-cell memorane protein) (CD48), inicial Homo sapiens CD47 antigen (Rh-related antigen, integrin-associated signal
NM_001///	transducer) (CD47), mRNA
NM_000733	Homo sapiens CD3E antigen, epsilon polypeptide (TiT3 complex) (CD3E), mRNA
NM 000732	Homo sapiens CD3D antigen, delta polypeptide (TiT3 complex) (CD3D),
_	mRNA
NM_001776	Homo sapiens ectonucleoside triphosphate diphosphohydrolase 1 (ENTPD1), mRNA
NM 001775	Homo sapiens CD38 antigen (p45) (CD38), mRNA
NM 001774	Homo sapiens CD37 antigen (CD37), mRNA
NM_001773	Homo sapiens CD34 antigen (CD34), mRNA  Homo sapiens CD34 antigen (CD34), mRNA
NM 003830	Homo sapiens CD34 antigen (CD34), intervi Homo sapiens sialic acid binding Ig-like lectin 5 (SIGLEC5), mRNA
NM 001245	Homo sapiens sialic acid binding Ig-like lectin 6 (SIGLEC6), mRNA
NM 001243	Homo sapiens CD33 antigen (gp67) (CD33), mRNA
NM_001772	Homo sapiens CD33 antigen (gp07) (CD33), fluctvi Homo sapiens CD2 antigen (p50), sheep red blood cell receptor (CD2), mRNA
	Homo sapiens CD22 antigen (CD22), mRNA
NM_001771	Homo sapiens CD12 antigen (CD22), hikvia  Homo sapiens CD1D antigen, d polypeptide (CD1D), mRNA
NM_001766	Homo sapiens CD1D antigen, a polypeptide (CD1D), indvA  Homo sapiens CD1C antigen, c polypeptide (CD1C), mRNA
NM_001765	nomo sapiens CDTC antigen, e porypeptide (CDTC), indian

	1 1 wide (CDID) mDNA
NM_001764	Homo sapiens CD1B antigen, b polypeptide (CD1B), mRNA
NM_001838	Homo sapiens chemokine (C-C motif) receptor 7 (CCR7), mRNA
NM_001837	Homo sapiens chemokine (C-C motif) receptor 3 (CCR3), mRNA
NM_001758	Homo sapiens cyclin D1 (PRAD1 parathyroid adenomatosis 1) (CCND1),
	mRNA (COVERD) PNA
NM_000731	Homo sapiens cholecystokinin B receptor (CCKBR), mRNA
NM_000730	Homo sapiens cholecystokinin A receptor (CCKAR), mRNA
NM_001757	Homo sapiens carbonyl reductase 1 (CBR1), mRNA
NM_001754	Homo sapiens runt-related transcription factor 1 (acute myeloid leukemia 1; aml1 oncogene) (RUNX1), mRNA
NM_003688	Homo sapiens calcium/calmodulin-dependent serine protein kınase (MAGUK
NM 001747	Homo sapiens capping protein (actin filament), gelsolin-like (CAPG), mRNA
NM_001744	Homo sapiens calcium/calmodulin-dependent protein kinase IV (CAMK4), mRNA
NM 001743	Homo sapiens calmodulin 2 (phosphorylase kinase, delta) (CALM2), mRNA
NM 001742	Homo saniens calcitonin receptor (CALCR), mRNA
NM 001741	Homo saniens calcitonin/calcitonin-related polypeptide, alpha (CALCA), mRNA
NM 000727	Homo sapiens calcium channel, voltage-dependent, gamma subunit 1
NW_000727	(CACNG1) mRNA
NM_000726	Homo sapiens calcium channel, voltage-dependent, beta 4 subunit (CACNB4),
NM_000725	Homo sapiens calcium channel, voltage-dependent, beta 3 subunit (CACNB3),
NM_000724	mRNA Homo sapiens calcium channel, voltage-dependent, beta 2 subunit (CACNB2), mRNA
NM_000723	Homo sapiens calcium channel, voltage-dependent, beta 1 subunit (CACNB1), mRNA
NM_000721	Homo sapiens calcium channel, voltage-dependent, alpha 1E subunit (CACNA1E), mRNA
NM_000720	Homo sapiens calcium channel, voltage-dependent, L type, alpha 1D subunit (CACNA1D), mRNA
NM_000719	Homo sapiens calcium channel, voltage-dependent, L type, alpha 1C subunit (CACNA1C), mRNA
NM_000718	Homo sapiens calcium channel, voltage-dependent, L type, alpha 1B subunit (CACNA1B), mRNA
NM_001739	Homo sapiens carbonic anhydrase VA, mitochondrial (CA5A), nuclear gene encoding mitochondrial protein, mRNA
ND ( 001729	Homo sapiens carbonic anhydrase I (CA1), mRNA
NM_001738	Homo sapiens complement component 9 (C9), mRNA
NM_001737	Homo sapiens complement component 5 receptor 1 (C5a ligand) (C5R1), mRNA
NM_001736	Homo sapiens complement component 5 (C5), mRNA
NM_001735	Homo sapiens compenent s (es), mad a Homo sapiens cholesterol 25-hydroxylase (CH25H), mRNA
NM 003956	Homo sapiens complement component 1, s subcomponent (C1S), mRNA
NM 001734	Homo sapiens complement component 1, r subcomponent (C1R), mRNA
NM_001733	Homo sapiens butyrophilin, subfamily 1, member A1 (BTN1A1), mRNA
NM_001732	Homo sapiens B-cell translocation gene 1, anti-proliferative (BTG1), mRNA
NM_001731	Homo sapiens betacellulin (BTC), mRNA
NM_001729	Homo sapiens betacentini (BTC), inclvid  Homo sapiens basigin (BSG), mRNA
NM_001728 NM_003742	Homo sapiens ATP-binding cassette, sub-family B (MDR/TAP), member 11
	(ABCB11), mRNA
NM_001727	Homo sapiens bombesin-like receptor 3 (BRS3), mRNA

NM 000059	Homo sapiens breast cancer 2, early onset (BRCA2), mRNA
NM 001725	Homo sapiens bactericidal/permeability-increasing protein (BPI), mRNA
NM 001724	Homo sapiens 2,3-bisphosphoglycerate mutase (BPGM), mRNA
NM 001723	Homo sapiens bullous pemphigoid antigen 1 (230/240kD) (BPAG1), mRNA
NM 001717	Homo sapiens basonuclin (BNC), mRNA
NM 001722	Homo sapiens BN51 (BHK21) temperature sensitivity complementing (BN51T),
1111_001/22	mRNA
NM 001721	Homo sapiens BMX non-receptor tyrosine kinase (BMX), mRNA
NM 001203	Homo sapiens bone morphogenetic protein receptor, type IB (BMPR1B), mRNA
NM 001720	Homo sapiens bone morphogenetic protein 8 (osteogenic protein 2) (BMP8),
11212_001720	mRNA
NM 001719	Homo sapiens bone morphogenetic protein 7 (osteogenic protein 1) (BMP7),
	mRNA
NM 001202	Homo sapiens bone morphogenetic protein 4 (BMP4), mRNA
NM 000713	Homo sapiens biliverdin reductase B (flavin reductase (NADPH)) (BLVRB),
	mRNA
NM_000712	Homo sapiens biliverdin reductase A (BLVRA), mRNA
NM 001713	Homo sapiens betaine-homocysteine methyltransferase (BHMT), mRNA
NM 001712	Homo sapiens carcinoembryonic antigen-related cell adhesion molecule 1
	(biliary glycoprotein) (CEACAM1), mRNA
NM 001711	Homo sapiens biglycan (BGN), mRNA
NM 000711	Homo sapiens bone gamma-carboxyglutamate (gla) protein (osteocalcin)
	(BGLAP), mRNA
NM 001709	Homo sapiens brain-derived neurotrophic factor (BDNF), mRNA
NM 000710	Homo sapiens bradykinin receptor B1 (BDKRB1), mRNA
NM 001707	Homo sapiens B-cell CLL/lymphoma 7B (BCL7B), mRNA
NM 001706	Homo sapiens B-cell CLL/lymphoma 6 (zinc finger protein 51) (BCL6), mRNA
NM 003921	Homo sapiens B-cell CLL/lymphoma 10 (BCL10), mRNA
NM 003657	Homo sapiens breast carcinoma amplified sequence 1 (BCAS1), mRNA
NM 001188	Homo sapiens BCL2-antagonist/killer 1 (BAK1), mRNA
NM 001704	Homo sapiens brain-specific angiogenesis inhibitor 3 (BAI3), mRNA
NM 001703	Homo sapiens brain-specific angiogenesis inhibitor 2 (BAI2), mRNA
NM 001702	Homo sapiens brain-specific angiogenesis inhibitor 1 (BAI1), mRNA
NM 001186	Homo sapiens BTB and CNC homology 1, basic leucine zipper transcription
_	factor 1 (BACH1), mRNA
NM 001701	Homo sapiens bile acid Coenzyme A amino acid N-acyltransferase (glycine N-
_	choloyltransferase) (BAAT), mRNA
NM_001185	Homo sapiens alpha-2-glycoprotein 1, zinc (AZGP1), mRNA
NM_001184	Homo sapiens ataxia telangiectasia and Rad3 related (ATR), mRNA
NM_000053	Homo sapiens ATPase, Cu++ transporting, beta polypeptide (Wilson disease)
_	(ATP7B), mRNA
NM_003945	Homo sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump) 9kD
-	(ATP6H), mRNA
NM_001696	Homo sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump)
_	31kD (ATP6E), mRNA
NM 001693	Homo sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump), beta
_	polypeptide, 56/58kD, isoform 2 (ATP6B2), mRNA
NM 001692	Homo sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump), beta
	polypeptide, 56/58kD, isoform 1 (ATP6B1), mRNA
NM_001691	Homo sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump),
	alpha polypeptide, 70kD, isoform 2 (ATP6A2), mRNA
NM 001690	Homo sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump),
[ 1ATAT 00 1 0 A O	none sapiens A11 ase, H+ transporting, tysosomai (vacuotai proton pump),

	alpha polypeptide, 70kD, isoform 1 (ATP6A1), mRNA
NM_001697	Homo sapiens ATP synthase, H+ transporting, mitochondrial F1 complex, O
_	subunit (oligomycin sensitivity conferring protein) (ATP50), mRNA
NM 001686	Homo sapiens ATP synthase, H+ transporting, mitochondrial F1 complex, beta
	polypeptide (ATP5B), nuclear gene encoding mitochondrial protein, mRNA
NM 000704	Homo sapiens ATPase, H+/K+ exchanging, alpha polypeptide (ATP4A), mRNA
NM 001684	Homo sapiens ATPase, Ca++ transporting, plasma membrane 4 (ATP2B4),
<b>-</b>	mRNA
NM_001682	Homo sapiens ATPase, Ca++ transporting, plasma membrane 1 (ATP2B1), mRNA
NM_001681	Homo sapiens ATPase, Ca++ transporting, cardiac muscle, slow twitch 2
	(ATP2A2), mRNA
NM_001679	Homo sapiens ATPase, Na+/K+ transporting, beta 3 polypeptide (ATP1B3),
	mRNA
NM_001678	Homo sapiens ATPase, Na+/K+ transporting, beta 2 polypeptide (ATP1B2), mRNA
NM_001677	Homo sapiens ATPase, Na+/K+ transporting, beta 1 polypeptide (ATP1B1), mRNA
NM 000703	Homo sapiens ATPase, Na+/K+ transporting, alpha 3 polypeptide (ATP1A3),
NML_000703	mRNA
NM_000702	Homo sapiens ATPase, Na+/K+ transporting, alpha 2 (+) polypeptide (ATP1A2), mRNA
NM 000701	Homo sapiens ATPase, Na+/K+ transporting, alpha 1 polypeptide (ATP1A1),
1441_000701	mRNA
NM_000051	Homo sapiens ataxia telangiectasia mutated (includes complementation groups
1111_00001	A, C and D) (ATM), mRNA
NM_001675	Homo sapiens activating transcription factor 4 (tax-responsive enhancer element B67) (ATF4), mRNA
NM 001673	Homo sapiens asparagine synthetase (ASNS), mRNA
NM 000048	Homo sapiens argininosuccinate lyase (ASL), mRNA
NM_001670	Homo sapiens armadillo repeat gene deletes in velocardiofacial syndrome
1411_001070	(ARVCF), mRNA
NM 001179	Homo sapiens ADP-ribosyltransferase 3 (ART3), mRNA
NM 000047	Homo sapiens arylsulfatase E (chondrodysplasia punctata 1) (ARSE), mRNA
NM_001178	Homo sapiens aryl hydrocarbon receptor nuclear translocator-like (ARNTL), mRNA
NM 001668	Homo sapiens aryl hydrocarbon receptor nuclear translocator (ARNT), mRNA
NM 001667	Homo sapiens ADP-ribosylation factor-like 2 (ARL2), mRNA
NM 001007	Homo sapiens Rho GDP dissociation inhibitor (GDI) gamma (ARHGDIG),
11117_001170	mRNA
NM 001665	Homo sapiens ras homolog gene family, member G (rho G) (ARHG), mRNA
NM 001661	Homo sapiens ADP-ribosylation factor 4-like (ARF4L), mRNA
NM 001659	Homo sapiens ADP-ribosylation factor 3 (ARF3), mRNA
NM 001657	Homo sapiens amphiregulin (schwannoma-derived growth factor) (AREG),
14141_001027	mRNA
NM_001654	Homo sapiens v-raf murine sarcoma 3611 viral oncogene homolog 1 (ARAF1),
1111_001054	mRNA
NM 001169	Homo sapiens aquaporin 8 (AQP8), mRNA
NM 001651	Homo sapiens aquaporin 5 (AQP5), mRNA
NM 001648	Homo sapiens addaporm's (1275), med 12  Homo sapiens kallikrein 3, (prostate specific antigen) (KLK3), mRNA
NM 000484	Homo sapiens amyloid beta (A4) precursor protein (protease nexin-II, Alzheimer
14141_000404	disease) (APP), mRNA
L	

<del></del>	TO A POD WARNA
NM_001647	Homo sapiens apolipoprotein D (APOD), mRNA
NM_001646	Homo sapiens apolipoprotein C-IV (APOC4), mRNA
NM_000384	Homo sapiens apolipoprotein B (including Ag(x) antigen) (APOB), mRNA
NM_001643	Homo sapiens apolipoprotein A-II (APOA2), mRNA
NM_001168	Homo sapiens baculoviral IAP repeat-containing 5 (survivin) (BIRC5), mRNA
NM_001167	Homo sapiens baculoviral IAP repeat-containing 4 (BIRC4), mRNA
NM_001164	Homo sapiens amyloid beta (A4) precursor protein-binding, family B, member 1 (Fe65) (APBB1), mRNA
NM_001163	Homo sapiens amyloid beta (A4) precursor protein-binding, family A, member 1 (X11) (APBA1), mRNA
NM_001161	Homo sapiens nudix (nucleoside diphosphate linked moiety X)-type motif 2 (NUDT2), mRNA
NM 001637	Homo sapiens acyloxyacyl hydrolase (neutrophil) (AOAH), mRNA
NM 001630	Homo sapiens annexin A8 (ANXA8), mRNA
NM 003568	Homo sapiens annexin A9 (ANXA9), mRNA
NM 000700	Homo sapiens annexin A1 (ANXA1), mRNA
	Homo sapiens solute carrier family 25 (mitochondrial carrier; adenine nucleotide
NM_001152	translocator), member 5 (SLC25A5), nuclear gene encoding mitochondrial protein, mRNA
NM_001151	Homo sapiens solute carrier family 25 (mitochondrial carrier; adenine nucleotide translocator), member 4 (SLC25A4), nuclear gene encoding mitochondrial protein, mRNA
NM_001150	Homo sapiens alanyl (membrane) aminopeptidase (aminopeptidase N,
14141_001130	aminopeptidase M, microsomal aminopeptidase, CD13, p150) (ANPEP), mRNA
NM 001146	Homo sapiens angiopoietin 1 (ANGPT1), mRNA
NM 000699	Homo sapiens amylase, alpha 2A; pancreatic (AMY2A), mRNA
NM 000481	Homo sapiens aminomethyltransferase (glycine cleavage system protein T)
14141_000461	(AMT), mRNA
NM_000480	Homo sapiens adenosine monophosphate deaminase (isoform E) (AMPD3), mRNA
NM 001144	Homo sapiens autocrine motility factor receptor (AMFR), mRNA
NM 001143	Homo sapiens amelogenin (Y chromosome) (AMELY), mRNA
NM 001633	Homo sapiens alpha-1-microglobulin/bikunin precursor (AMBP), mRNA
NM 000698	Homo sapiens arachidonate 5-lipoxygenase (ALOX5), mRNA
	Homo sapiens arachidonate 15-lipoxygenase (ALOX15), mRNA
NM_001140	Homo sapiens arachidonate 12-lipoxygenase, 12R type (ALOX12B), mRNA
NM_001139	Homo sapiens arachidanete 12 inoverses (ALOVI2) mPNA
NM_000697	Homo sapiens arachidonate 12-lipoxygenase (ALOX12), mRNA
NM_001628	Homo sapiens aldo-keto reductase family 1, member B1 (aldose reductase) (AKR1B1), mRNA
NM_000696	Homo sapiens aldehyde dehydrogenase 9 (gamma-aminobutyraldehyde dehydrogenase, E3 isozyme) (ALDH9), mRNA
NM_000692	Homo sapiens aldehyde dehydrogenase 5 (ALDH5), mRNA
NM 003748	Homo sapiens aldehyde dehydrogenase 4 (glutamate gamma-semialdehyde
	dehydrogenase; pyrroline-5-carboxylate dehydrogenase) (ALDH4), mRNA
NM 000690	Homo sapiens aldehyde dehydrogenase 2, mitochondrial (ALDH2), mRNA
NM 000689	Homo sapiens aldehyde dehydrogenase 1, soluble (ALDH1), mRNA
NM 001627	Homo sapiens activated leucocyte cell adhesion molecule (ALCAM), mRNA
NM_000688	Homo sapiens aminolevulinate, delta-, synthase 1 (ALAS1), nuclear gene
14141_000000	encoding mitochondrial protein, mRNA
NIM 002690	Homo sapiens aldo-keto reductase family 7, member A2 (aflatoxin aldehyde
NM_003689	reductase) (AKR7A2), mRNA
NM 003886	Homo sapiens A kinase (PRKA) anchor protein 4 (AKAP4), mRNA

NM 003488	Homo sapiens A kinase (PRKA) anchor protein 1 (AKAP1), mRNA
NM 001622	Homo sapiens alpha-2-HS-glycoprotein (AHSG), mRNA
NM 003659	Homo sapiens alkylglycerone phosphate synthase (AGPS), mRNA
NM 001133	Homo sapiens afamin (AFM), mRNA
NM 001131	Homo sapiens acidic epididymal glycoprotein-like 1 (AEGL1), mRNA
NM_003938	Homo sapiens adaptor-related protein complex 3, delta 1 subunit (AP3D1),
	mRNA
NM_001127	Homo sapiens adaptor-related protein complex 1, beta 1 subunit (AP1B1),
	mRNA
NM 000676	Homo sapiens adenosine A2b receptor (ADORA2B), mRNA
NM 000674	Homo sapiens adenosine A1 receptor (ADORA1), mRNA
NM 001124	Homo sapiens adrenomedullin (ADM), mRNA
NM 001120	Homo sapiens tetracycline transporter-like protein (TETRAN), mRNA
NM_001118	Homo sapiens adenylate cyclase activating polypeptide 1 (pituitary) receptor
<del></del>	type I (ADCYAP1R1), mRNA
NM 000666	Homo sapiens aminoacylase 1 (ACY1), mRNA
NM 001613	Homo sapiens actin, alpha 2, smooth muscle, aorta (ACTA2), mRNA
NM 001097	Homo sapiens acrosin (ACR), mRNA
NM 003501	Homo sapiens acyl-Coenzyme A oxidase 3, pristanoyl (ACOX3), mRNA
NM 003500	Homo sapiens acyl-Coenzyme A oxidase 2, branched chain (ACOX2), mRNA
NM 001098	Homo sapiens aconitase 2, mitochondrial (ACO2), nuclear gene encoding
14141_001030	mitochondrial protein, mRNA
NM 001096	Homo sapiens ATP citrate lyase (ACLY), mRNA
NM_001609	Homo sapiens acyl-Coenzyme A dehydrogenase, short/branched chain
1111_001003	(ACADSB), nuclear gene encoding mitochondrial protein, mRNA
NM 001608	Homo sapiens acyl-Coenzyme A dehydrogenase, long chain (ACADL), mRNA
NM 001093	Homo sapiens acetyl-Coenzyme A carboxylase beta (ACACB), mRNA
NM 001089	Homo sapiens ATP-binding cassette, sub-family A (ABC1), member 3
112.2_00.000	(ABCA3), mRNA
NM 000663	Homo sapiens 4-aminobutyrate aminotransferase (ABAT), nuclear gene
	encoding mitochondrial protein, mRNA
NM 001605	Homo sapiens alanyl-tRNA synthetase (AARS), mRNA
NM 021123	Homo sapiens G antigen 7 (GAGE7), mRNA
NM 006994	Homo sapiens butyrophilin, subfamily 3, member A3 (BTN3A3), mRNA
NM 001812	Homo sapiens centromere protein C 1 (CENPC1), mRNA
NM_015983	Homo sapiens ubiquitin-conjugating enzyme HBUCE1 (LOC51619), mRNA
NM 009590	Homo sapiens amine oxidase, copper containing 2 (retina-specific) (AOC2),
_	transcript variant 2, mRNA
NM 001159	Homo sapiens aldehyde oxidase 1 (AOX1), mRNA
NM 007326	Homo sapiens diaphorase (NADH) (cytochrome b-5 reductase) (DIA1), nuclear
	gene encoding mitochondrial protein, transcript variant S, mRNA
NM_005158	Homo sapiens v-abl Abelson murine leukemia viral oncogene homolog 2 (arg,
_	Abelson-related gene) (ABL2), transcript variant a, mRNA
NM_004441	Homo sapiens EphB1 (EPHB1) mRNA
NM 004089	Homo sapiens delta sleep inducing peptide, immunoreactor (DSIPI), mRNA
NM 004077	Homo sapiens citrate synthase (CS), nuclear gene encoding mitochondrial
_	protein, mRNA
NM 003890	Homo sapiens IgG Fc binding protein (FC(GAMMA)BP) mRNA
NM 003582	Homo sapiens dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 3
_	(DYRK3) mRNA
NM_001396	Homo sapiens dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 1
_	(DYRK1) mRNA

### **CLAIMS**

#### What we claim is:

5

15

20

25

1. A double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene, wherein said siNA molecule comprises one or more chemical modifications and each strand of said double-stranded siNA comprises about 21 nucleotides.

- 2. The siNA molecule of claim 1, wherein said siNA molecule comprises no ribonucleotides.
- 3. The siNA molecule of claim 1, wherein said siNA molecule comprises ribonucleotides.
  - 4. The siNA molecule of claim 1, wherein one of the strands of said double-stranded siNA molecule comprises a nucleotide sequence that is complementary to a nucleotide sequence of the endogenous mammalian target gene or a portion thereof, and wherein the second strand of said double-stranded siNA molecule comprises a nucleotide sequence substantially similar to the nucleotide sequence of the endogenous mammalian target gene or a portion thereof.
  - 5. The siNA molecule of claim 4, wherein each strand of the siNA molecule comprises about 19 to about 23 nucleotides, and wherein each strand comprises at least about 19 nucleotides that are complementary to the nucleotides of the other strand.
  - 6. The siNA molecule of claim 1, wherein said siNA molecule comprises an antisense region comprising a nucleotide sequence that is complementary to a nucleotide sequence of the endogenous mammalian target gene or a portion thereof, and wherein said siNA further comprises a sense region, wherein said sense region comprises a nucleotide sequence substantially similar to the nucleotide sequence of said endogenous mammalian target gene or a portion thereof.
- 7. The siNA molecule of claim 6, wherein said antisense region and said sense region each comprise about 19 to about 23 nucleotides, and wherein said antisense region comprises at least about 19 nucleotides that are complementary to nucleotides of the sense region.

8. The siNA molecule of claim 1, wherein said siNA molecule comprises a sense region and an antisense region and wherein said antisense region comprises a nucleotide sequence that is complementary to a nucleotide sequence of RNA encoded by the endogenous mammalian target gene or a portion thereof and said sense region comprises a nucleotide sequence that is complementary to said antisense region.

- 9. The siNA molecule of claim 6, wherein said siNA molecule is assembled from two separate oligonucleotide fragments, wherein one fragment comprises the sense region and the second fragment comprises the antisense region of said siNA molecule.
- 10. The siNA molecule of claim claim 6, wherein said sense region is connected to the antisense region via a linker molecule.
- 11. The siNA molecule of claim 10, wherein said linker molecule is a polynucleotide linker.
- 15 12. The siNA molecule of claim 10, wherein said linker molecule is a non-nucleotide linker.
  - 13. The siNA molecule of claim 6, wherein pyrimidine nucleotides in the sense region are 2'-O-methyl pyrimidine nucleotides.
- 14. The siNA molecule of claim 6, wherein purine nucleotides in the sense region are 2'-deoxy purine nucleotides.
  - 15. The siNA molecule of claim 6, wherein the pyrimidine nucleotides present in the sense region are 2'-deoxy-2'-fluoro pyrimidine nucleotides.
- The siNA molecule of claim 9, wherein the fragment comprising said sense region includes a terminal cap moiety at the 5'-end, the 3'-end, or both of the 5' and 3' ends of the fragment comprising said sense region.
  - 17. The siNA molecule of claim 16, wherein said terminal cap moiety is an inverted deoxy abasic moiety.
  - 18. The siNA molecule of claim 6, wherein the pyrimidine nucleotides of said antisense region are 2'-deoxy-2'-fluoro pyrimidine nucleotides.

5

19. The siNA molecule of claim 6, wherein the purine nucleotides of said antisense region are 2'-O-methyl purine nucleotides.

- 20. The siNA molecule of claim 6, wherein the purine nucleotides present in said antisense region comprise 2'-deoxy- purine nucleotides.
- 5 21. The siNA molecule of claim 18, wherein said antisense region comprises a phosphorothioate internucleotide linkage at the 3' end of said antisense region.
  - 22. The siNA molecule of claim 6, wherein said antisense region comprises a glyceryl modification at the 3' end of said antisense region.
- The siNA molecule of claim 9, wherein each of the two fragments of said siNA molecule comprise 21 nucleotides.
  - 24. The siNA molecule of claim 23, wherein about 19 nucleotides of each fragment of the siNA molecule are base-paired to the complementary nucleotides of the other fragment of the siNA molecule and wherein at least two 3' terminal nucleotides of each fragment of the siNA molecule are not base-paired to the nucleotides of the other fragment of the siNA molecule.
  - 25. The siNA molecule of claim 24, wherein each of the two 3' terminal nucleotides of each fragment of the siNA molecule are 2'-deoxy-pyrimidines.
  - 26. The siNA molecule of claim 25, wherein said 2'-deoxy-pyrimidine is 2'-deoxy-thymidine.
- 20 27. The siNA molecule of claim 23, wherein all 21 nucleotides of each fragment of the siNA molecule are base-paired to the complementary nucleotides of the other fragment of the siNA molecule.
  - 28. The siNA molecule of claim 23, wherein about 19 nucleotides of the antisense region are base-paired to the nucleotide sequence of the RNA encoded by the endogenous mammalian target gene or a portion thereof.
    - 29. The siNA molecule of claim 23, wherein 21 nucleotides of the antisense region are base-paired to the nucleotide sequence of the RNA encoded by the endogenous mammalian target gene or a portion thereof.
- 30. The siNA molecule of claim 9, wherein the 5'-end of the fragment comprising said antisense region optionally includes a phosphate group.

15

31. The siNA molecule of claim 1, wherein said mammalian gene is a human gene.

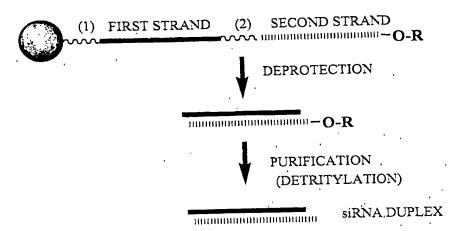
- 32. A double-stranded short interfering nucleic acid (siNA) molecule that inhibits the expression of an endogenous mammalian target RNA sequence, wherein each strand of said double-stranded siNA molecule comprises about 21 nucleotides and wherein said siNA molecule comprises no ribonucleotides.
- 33. The siNA molecule of claim 32, wherein said target RNA sequence is encoded by a human gene.
- A double-stranded short interfering nucleic acid (siNA) molecule that inhibits the expression of an endogenous mammalian target gene, wherein each strand of said double-stranded siNA molecule comprises about 21 nucleotides and wherein said siNA molecule does not require the presence of a ribonucleotide within the siNA molecule for the inhibition of expression of an endogenous mammalian target gene.
- The siNA molecule of claim 34, wherein said mammalian target gene is a human gene.
  - 36. The siNA molecule of claim 31 or claim 35, wherein said human gene is vascular endothelial growth factor (VEGF).
  - 37. The siNA molecule of claim 31 or claim 35, wherein said human gene is a receptor for VEGF.
- 20 38. The siNA of claim 37, wherein said receptor is VEGFR1.
  - 39. The siNA of claim 37, wherein said receptor is VEGFR2.
  - 40. The siNA of claim 37, wherein said receptor is VEGFR3
  - 41. The siNA molecule of claim 31 or claim 35, wherein said human gene is BCL2.
- 42. The siNA molecule of claim 31 or claim 35, wherein said human gene is HER2/neu.
  - 43. The siNA molecule of claim 31 or claim 35, wherein said human gene is c-Myc.
  - 44. The siNA molecule of claim 31 or claim 35, wherein said human gene is PCNA.
  - 45. The siNA molecule of claim 31 or claim 35, wherein said human gene is REL-A.

46. The siNA molecule of claim 31 or claim 35, wherein said human gene is PTP1B.

- 47. The siNA molecule of claim 31 or claim 35, wherein said human gene is BACE.
- 48. The siNA molecule of claim 31 or claim 35, wherein said human gene is CHK1.
- 49. The siNA molecule of claim 31 or claim 35, wherein said human gene is PKC-5 alpha.
  - 50. The siNA molecule of claim 31 or claim 35, wherein said human gene is EGFR (HER1).
  - 51. A pharmaceutical composition comprising the siNA molecule of claim 1 in an acceptable carrier or diluent.
- 10 52. Medicament comprising the siNA molecule of claim 1.
  - 53. Active ingredient comprising the siNA molecule of claim 1.
  - 54. Use of a double-stranded short interfering nucleic acid (siNA) molecule to down-regulate expression of an endogenous mammalian target gene, wherein said siNA molecule comprises one or more chemical modifications and each strand of said double-stranded siNA comprises about 21 nucleotides.

1/34

## Figure 1



= SOLID SUPPORT

R = TERMINAL PROTECTING GROUP FOR EXAMPLE: DIMETHOXYTRITYL (DMT)

= CLEAVABLE LINKER
(FOR EXAMPLE: NUCLEOTIDE SUCCINATE OR
(2) INVERTED DEOXYABASIC SUCCINATE)

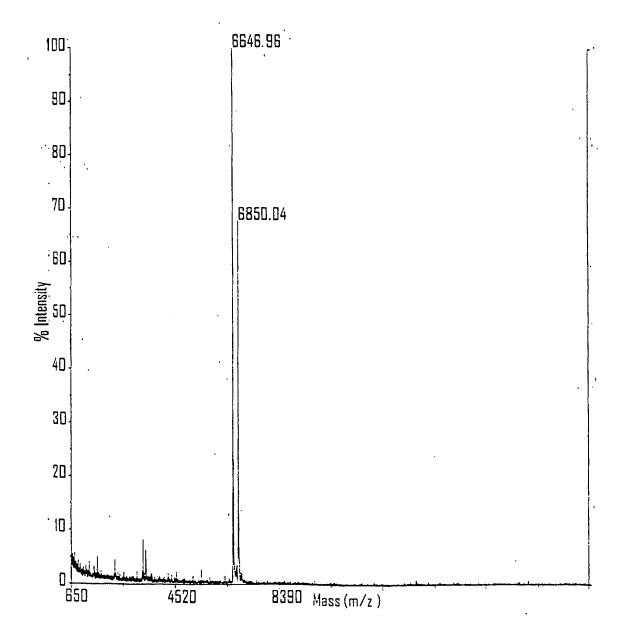
(FOR EXAMPLE: NUCLEOTIDE SUCCINATE OR INVERTED DEOXYABASIC SUCCINATE)

INVERTED DEOXYABASIC SUCCINATE LINKAGE

GLYCERYL SUCCINATE LINKAGE

2/34

## Figure 2



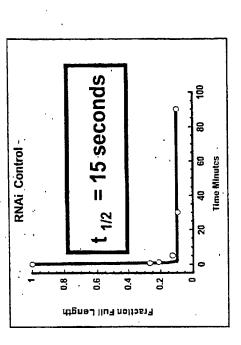
# Figure 3

5'-CGUACGCGGAAUACUUCGATT (SEQ ID NO: 925) 3'-TTGCAUGCGCCUUAUGAAGCU (SEQ ID NO: 926)

T % = 138 min5'-B cAAccAcAAAuAcAAcAATT B (SEQ ID NO: 925) 3'-TXGuuGGuGuuuuAuGuuGuu (SEQ ID NO: 927)

5'-B cAAccACAAAUAcAACAATT B (SEQ ID NO: 925) T 1/2 = 3.7 days 3'-TDGuuGGuGuuuuAuGuuGuu (SEQ ID NO: 928) 5'-B cAAccACAAAUAcAATT B (SEQ ID NO: 925 ) T % = 72 minutes 3'-XTGuuGGuGuuuuAuGuuGuu (SEQ ID NO: 929 )

5'-B cAAccACAAAUACAACAATT B (SEQ ID NO; 925) T 1/2 = 40 days 3'-LTGuuGGuGuuuuAuGuuGuu (SEQ ID NO: 930) 5'-B cAAccACAAAAUACAACAATT B (SEQ ID NO: 925) T 1/2 = 32 days 3'-tTGuuGGuGuuuuAuGuuGuu (SEQ ID NO: 931)



G, A, U, C = Guanosine, Adenosine, Uridine, Cytidine T = Thymidine

= phosphorothioate

Lower Case = 2'-deoxy-2'-fluoro

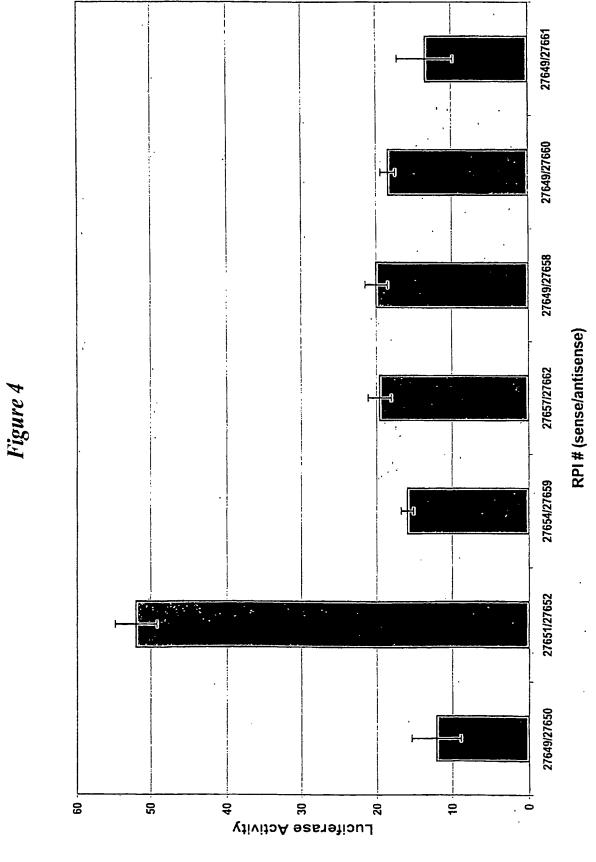
B = inverted deoxyabasic

D = inverted Thymidine G = terminal glycine

= Glyceryl moiety t = L-thymidine

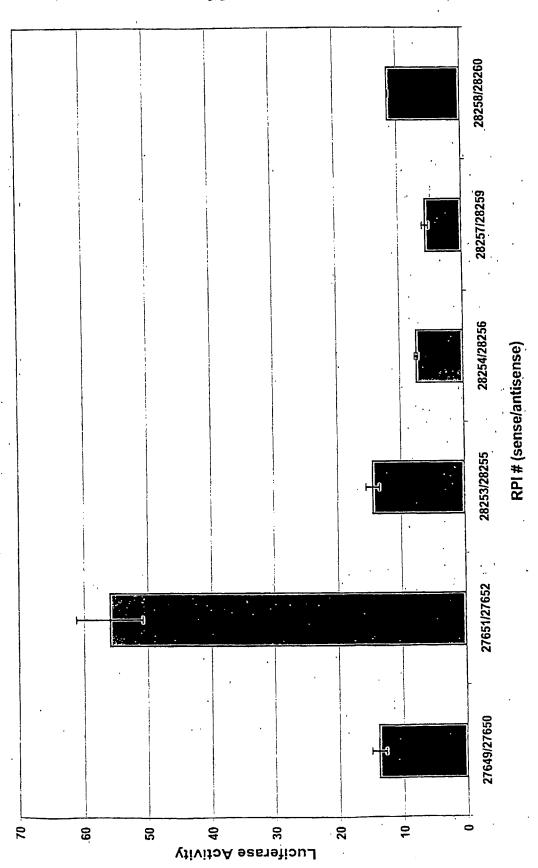
X = 3'-deoxy Thymidine

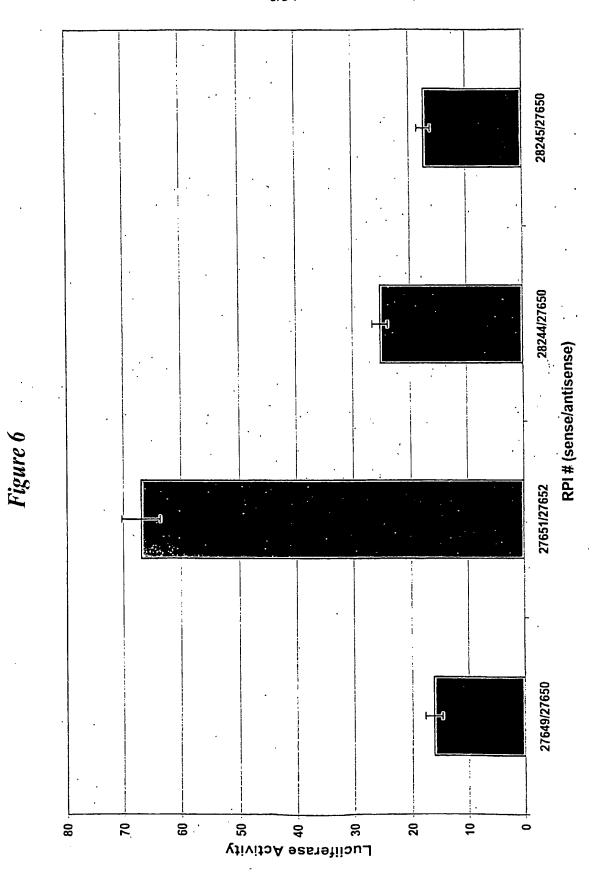
SUBSTITUTE SHEET (RULE 26)



SUBSTITUTE SHEET (RULE 26)

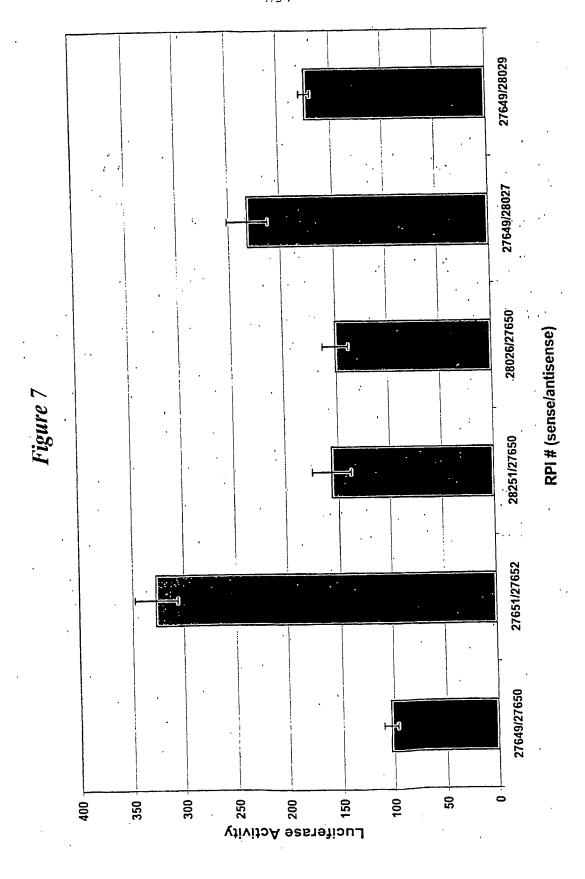




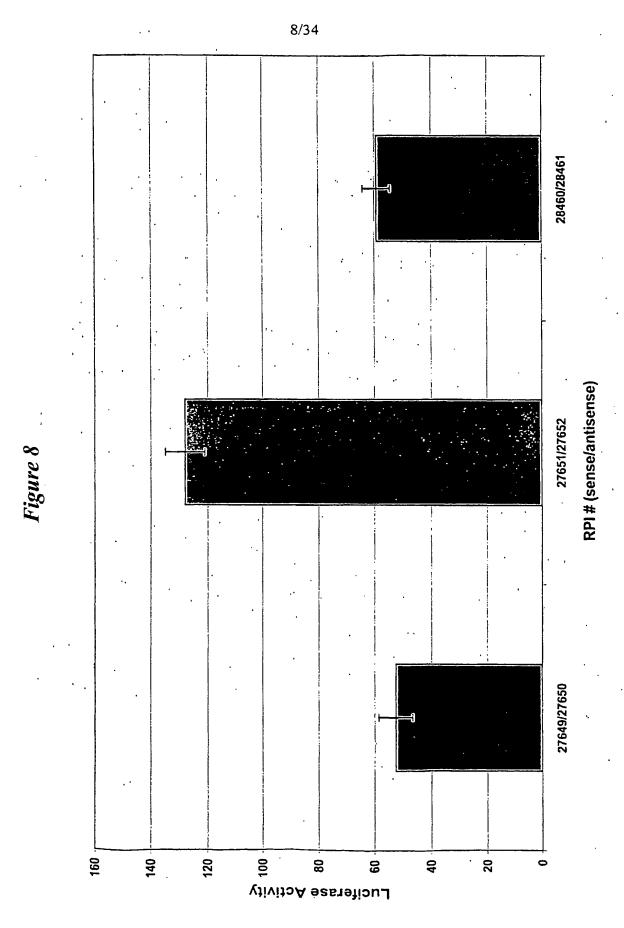


SUBSTITUTE SHEET (RULE 26)

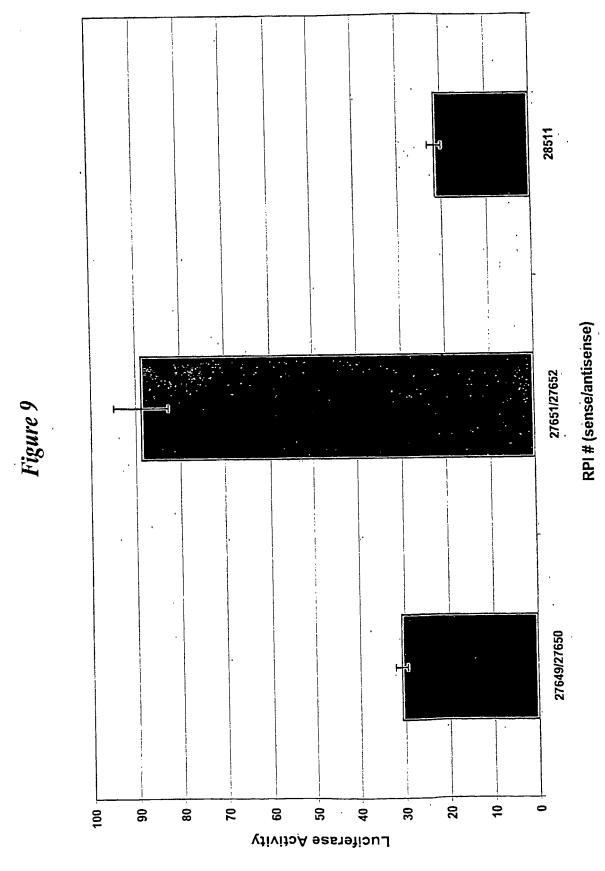




## SUBSTITUTE SHEET (RULE 26)

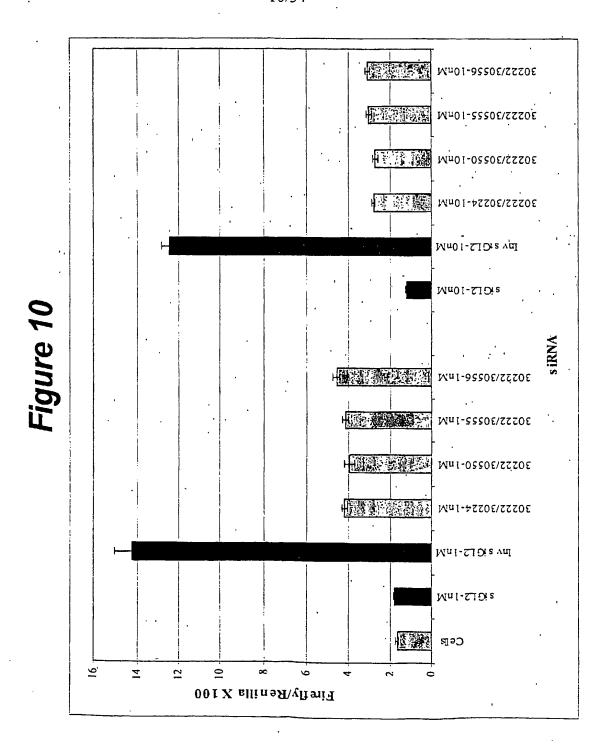


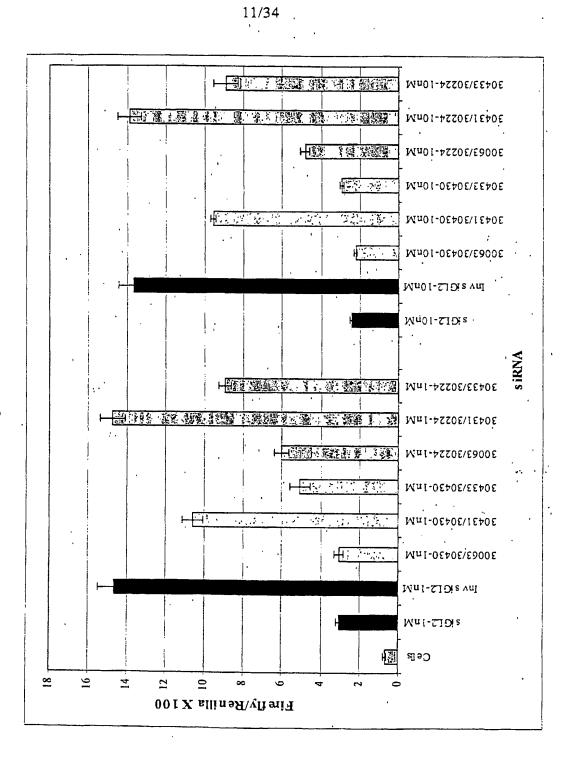
SUBSTITUTE SHEET (RULE 26)



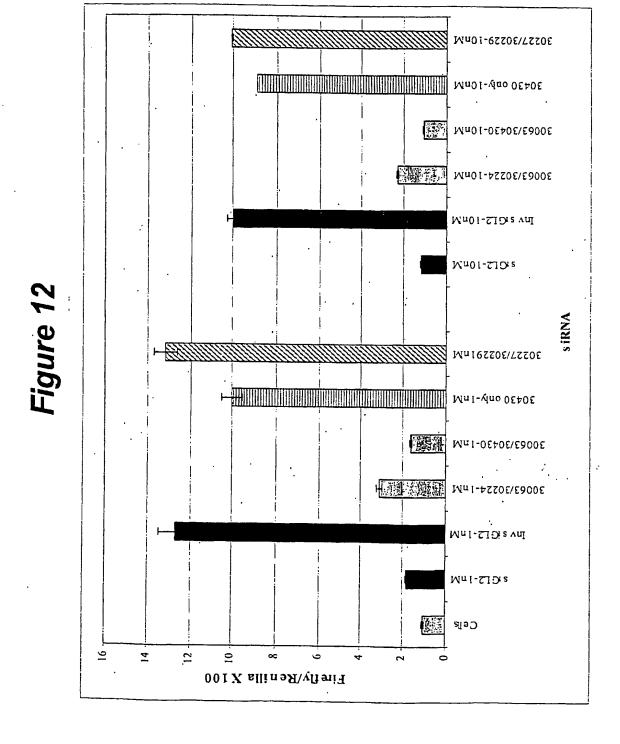
SUBSTITUTE SHEET (RULE 26)

10/34



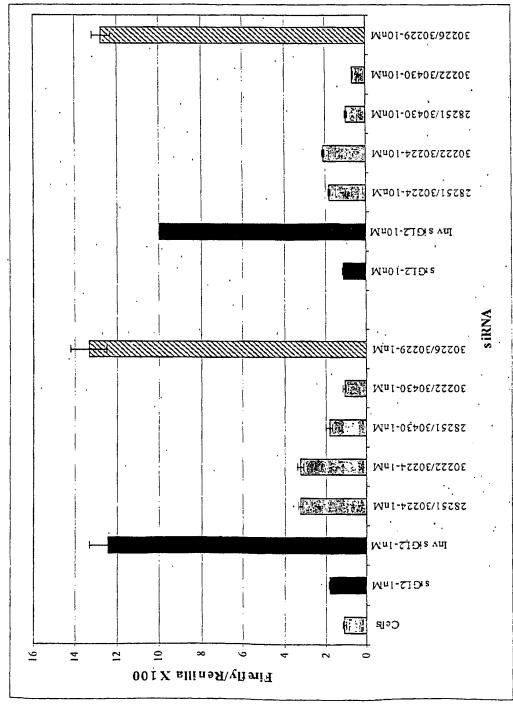


**SUBSTITUTE SHEET (RULE 26)** 



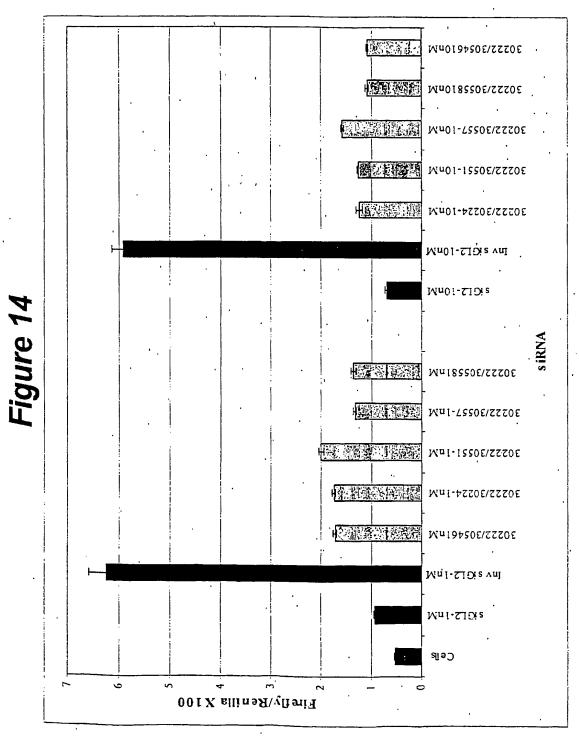
SUBSTITUTE SHEET (RULE 26)



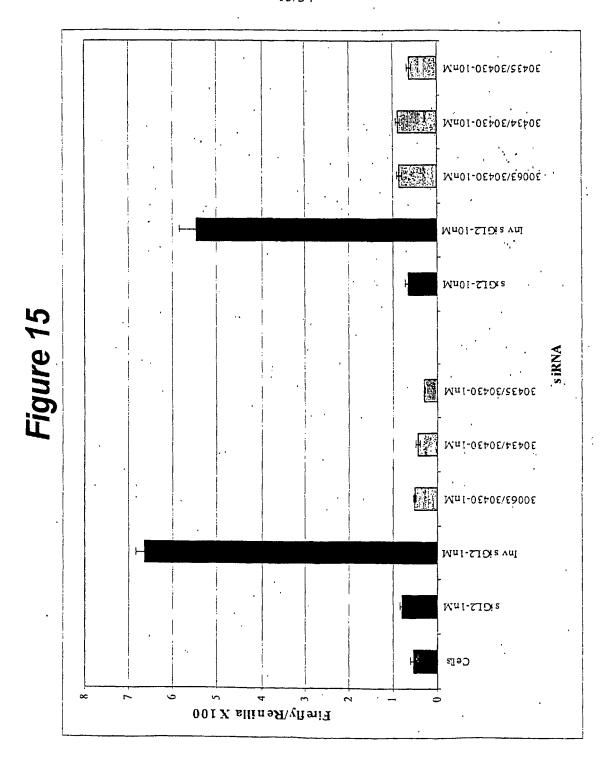


13/34

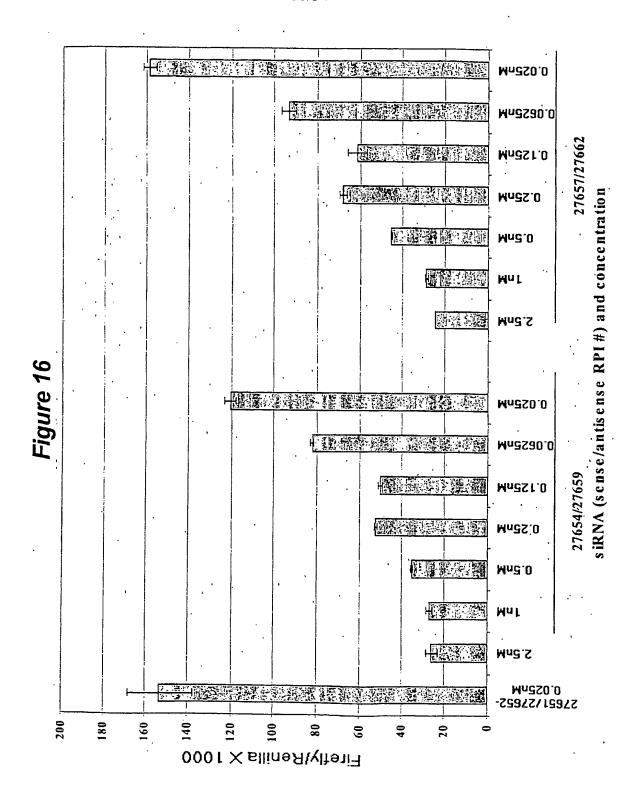




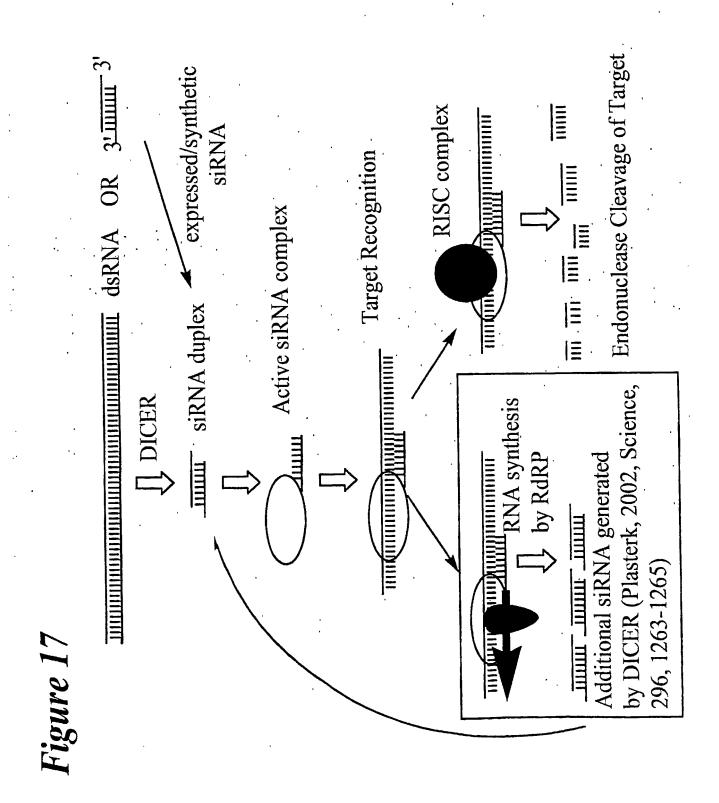




16/34



SUBSTITUTE SHEET (RULE 26)



# Figure 18

```
SENSE STRAND (SEQ ID NO 903)
              ALL PYRIMIDINES = 2'-O-ME OR 2'-FLUORO EXCEPT POSITIONS (N N)
       5'-
                -3'
       3'-
           L-(N_sN)NNNNNNNNNNNNNNNNNSN_sN_sN_sN_s
                                                             -5'
                          ANTISENSE STRAND (SEQ ID NO 904)
                   ALL PYRIMIDINES = 2'-FLUORO EXCEPT POSITIONS (N N)
                          SENSE STRAND (SEQ ID NO 905)
              ALL PYRIMIDINES = 2'-O-ME OR 2'-FLUORO EXCEPT POSITIONS (N N)
       5'-
                '-3'
{f B}
            L-(NN) NNNNNNNNNNNNNNNNNNNN
                                                             -5'
                          ANTISENSE STRAND (SEQ ID NO 906)
                    ALL PYRIMIDINES = 2'-FLUORO EXCEPT POSITIONS (N;N)
                         SENSE STRAND (SEQ ID NO 907)
              ALL PYRIMIDINES = 2'-O-ME OR 2'-FLUORO EXCEPT POSITIONS (N N)
                -3'
       3'-
             -5
                          ANTISENSE STRAND (SEQ ID NO 908)
                    ALL PYRIMIDINES = 2'-FLUORO EXCEPT POSITIONS (N N)
                        SENSE STRAND (SEQ ID NO 909)
      ALL PYRIMIDINES = 2'-FLUORO EXCEPT POSITIONS (N N) AND ALL PURINES = 2'-DEOXY
      5'-
                -3'
      3'-
           L-(N<sub>1</sub>N) NNNNNNNNNNNNNNNNNN
                                                             -5'
                       ANTISENSE STRAND (SEQ ID NO 910)
       ALL'PYRIMIDINES = 2'-FLUORO AND ALL PURINES = 2'-O-ME EXCEPT POSITIONS (N N)
                          SENSE STRAND (SEQ ID NO 911)
                  ALL PYRIMIDINES = 2'-FLUORO EXCEPT POSITIONS (N N)
      5'-
                -3'
\mathbf{E}
                                                             -5'
          L-(NN) NNNNNNNNNNNNNNNNNNNN
                       ANTISENSE STRAND (SEQ ID NO 912)
       ALL PYRIMIDINES = 2'-FLUORO AND ALL PURINES \approx 2'-O-ME EXCEPT POSITIONS (N N)

    SENSE STRAND (SEQ ID NO 909)

      ALL PYRIMIDINES = 2'-FLUORO EXCEPT POSITIONS (N N) AND ALL PURINES = 2'-DEOXY
      5'-
               -3'
F
      3'-
           -5'
                       ANTISENSE STRAND (SEQ ID NO 913)
       LL PYRIMIDINES = 2'-FLUORO EXCEPT POSITIONS (N N) AND ALL PURINES = 2'-DEOXY
```

POSITIONS (NN) CAN COMPRISE ANY NUCLEOTIDE, SUCH AS DEOXYNUCLEOTIDES (eg. THYMIDINE) OR UNIVERSAL BASES

B = ABASIC, INVERTED ABASIC, INVERTED NUCLEOTIDE OR OTHER TERMINAL CAP THAT IS OPTIONALLY PRESENT

L = GLYCERYL MOIETY THAT IS OPTIONALLY PRESENT

S = PHOSPHOROTHIOATE OR PHOSPHORODITHIOATE

Fig	ure 19	·	·
		SENSE STRAND (SEQ ID NO 914)	
A	5'- 3'-	$u_Su_Su_SG_Su$ c u u c c A u u c c A $u_Su_SG_ST_ST$ L- $T_ST$ A A A c A G A A G G u A A G $G_Su_SA_SA_Sc$ ANTISENSE STRAND (SEQ ID NO 915)	-3' -5'
	( .	SENSE STRAND (SEQ ID NO 916)	
В	5'- 3'-	uuuGucuuccAuuccAuuGTT L-TTAAAcAGAAGGuAAC ANTISENSE STRAND (SEQ ID NO 917)	-3' -5'
		SENSE STRAND (SEQ ID NO 918)	
<b>C</b>	5'-	iB-uuuGucuuccAuuccAuuGTT-iB L-T <sub>S</sub> TAAAcAGAAGGuAAGGuAAc ANTISENSE STRAND (SEQ ID NO 919)	-3' -5'
,		SENSE STRAND (SEQ ID NO 920)	
D	5'- 3'-	iB-uuuGucuuccAuuccAuuGTT-iB L-T <sub>S</sub> Taaacagaagguaagguaac ANTISENSE STRAND (SEQ ID NO 921).	-3' -5'
		SENSE STRAND (SEQ ID NO 922)	. }
<b>E</b>	5'- 3'-	iB-uuuGucuuccAuuccAuuGTT-iB L-TTaaacagaagguaagguaac ANTISENSE STRAND (SEQ ID NO 923)	-3' -5'
		SENSE STRAND (SEO ID NO 620)	· J
F	5'-	iB-uuu GucuuccAuuccAuuGTT-iB L-T <sub>S</sub> TAAAcAGAAGGuAAGGuAAC ANTISENSE STRAND (SEQ ID NO 924)	-3' -5'
	_		

lower case = 2'-O-Methyl or 2'-deoxy-2'-fluoro; italic lower case = 2'-deoxy-2'-fluoro ITALIC UPPER CASE = DEOXY

B = INVERTED DEOXYABASIC L = GLYCERYL MOIETY OPTIONALLY PRESENT S = PHOSPHOROTHIOATE OR PHOSPHORODITHIOATE

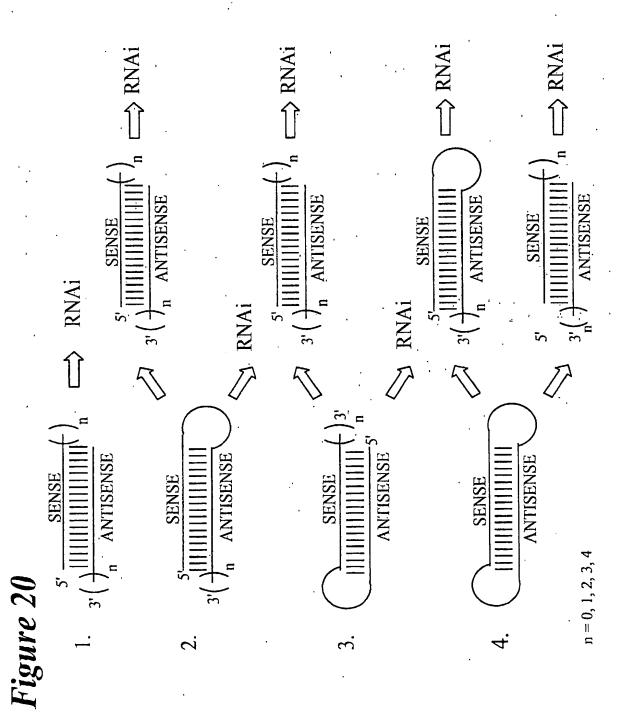
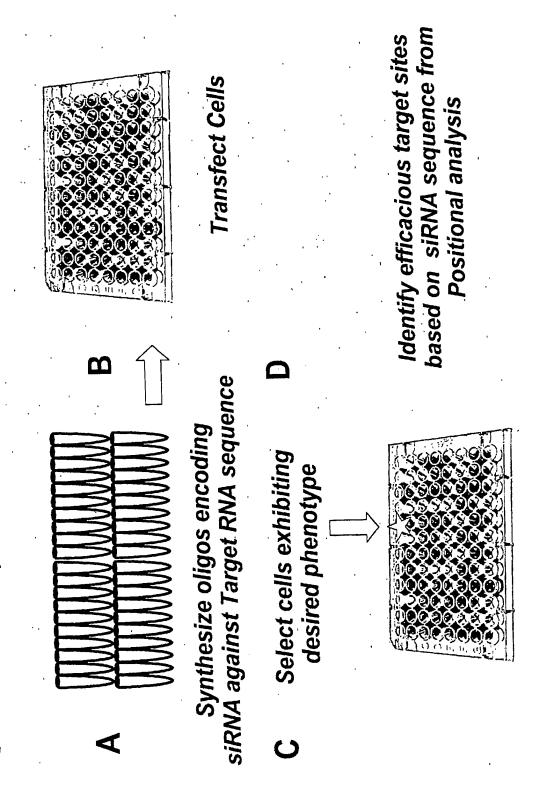
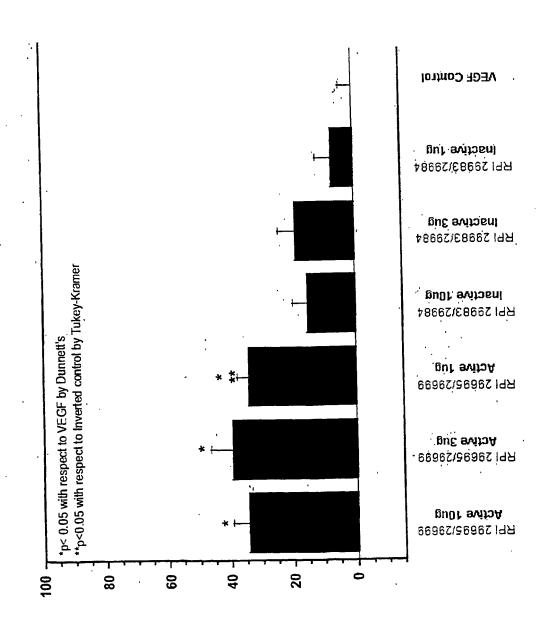


Figure 21: Target site Selection using siRNA



R = O, S, N, alkyl, substituted alkyl, O-alkyl, S-alkyl, alkaryl, or aralkyl B = Independently any nucleotide base, either naturally occurring or chemically modified, or optionally H (abasic).

Figure 23: Inhibition of VEGF-Induced Angiogenesis by siRNAs



% Inhibition of VEGF induced % sizenegoignA

# Figure 24: Modification Strategy

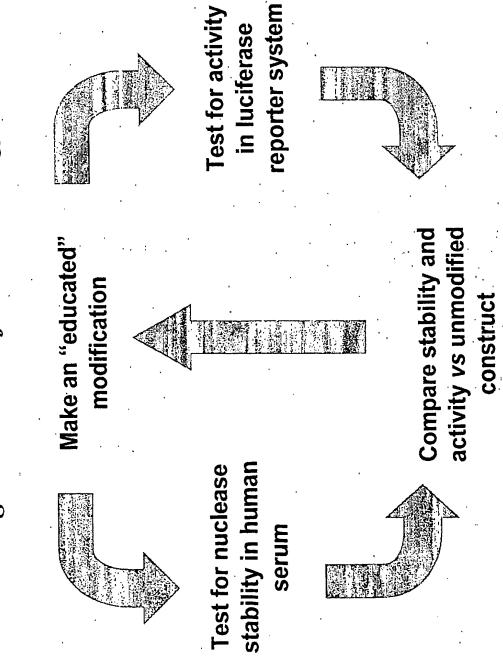


Figure 25: A549 24h EGFR (HER1) mRNA Expression

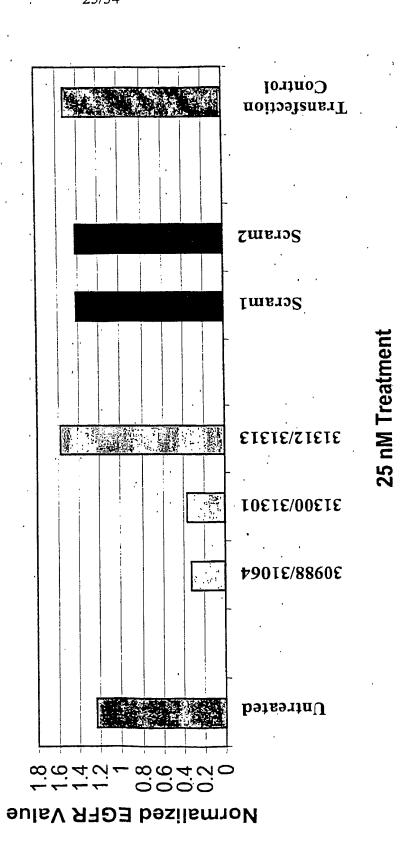


Figure 26: A549 24h PKCa mRNA Expression

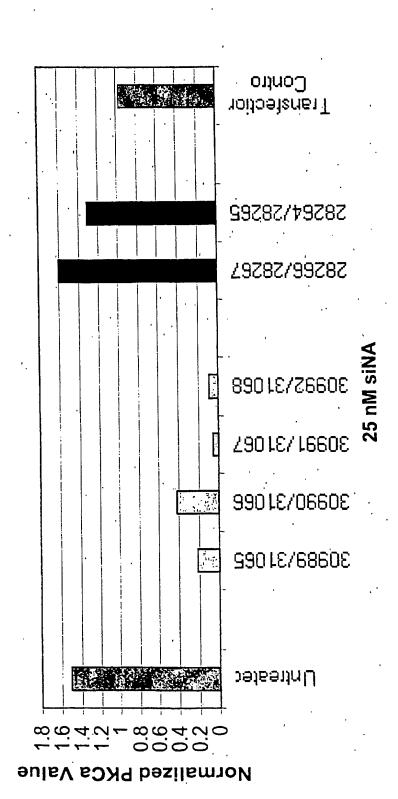


Figure 27: siNA mediated inhibition of MYC RNA

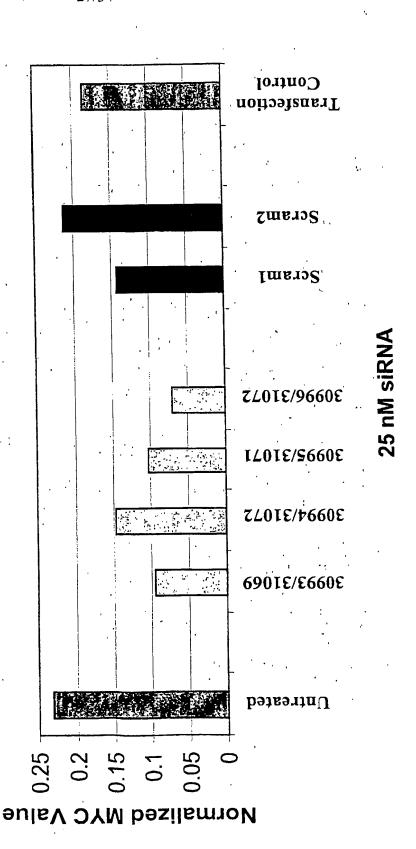


Figure 28: A549 24h Bcl2 mRNA Expression Screen

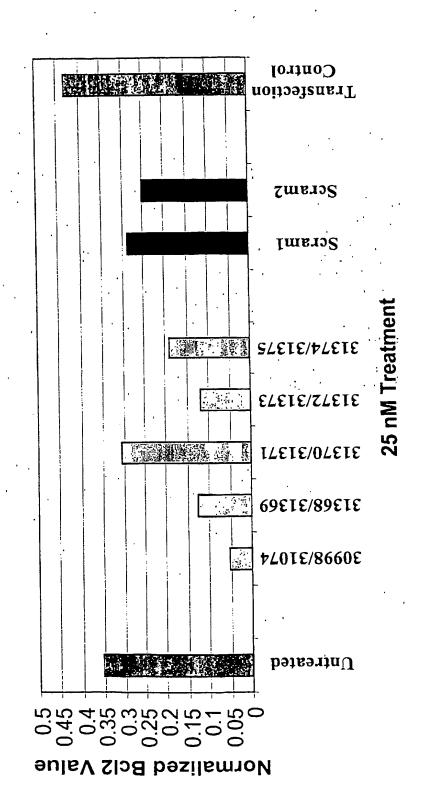
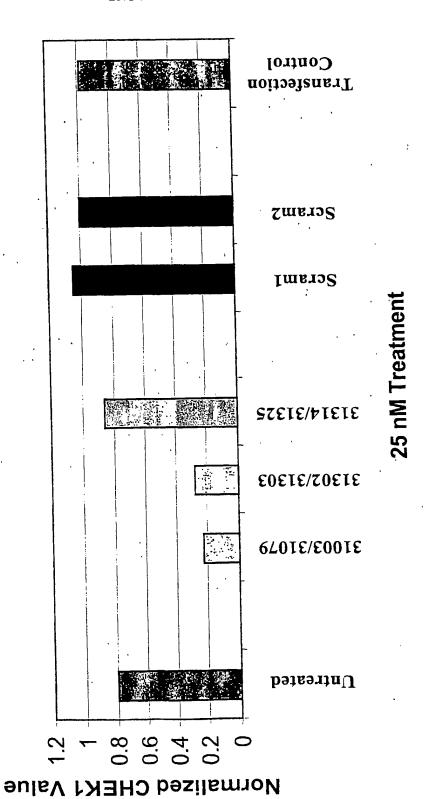


Figure 29: A549 24h CHEK1 mRNA Expression



30/34

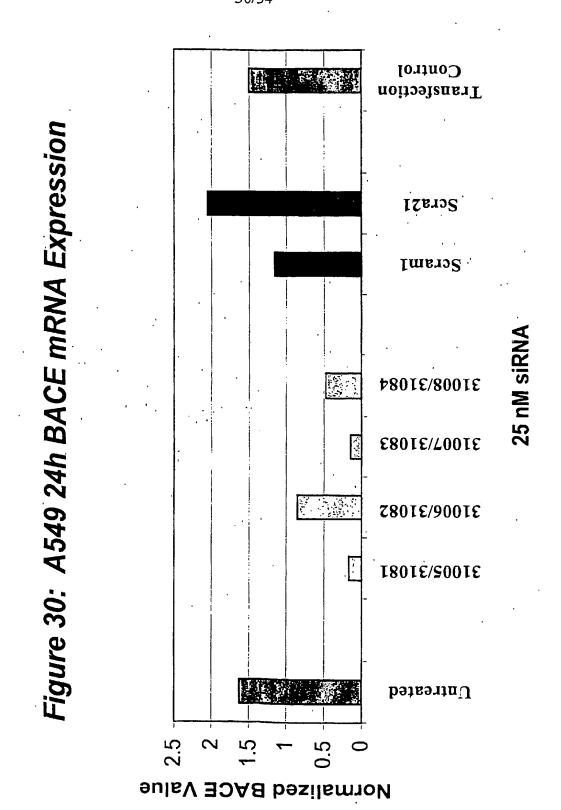


Figure 31: A549 24h CCND1 mRNA Expression

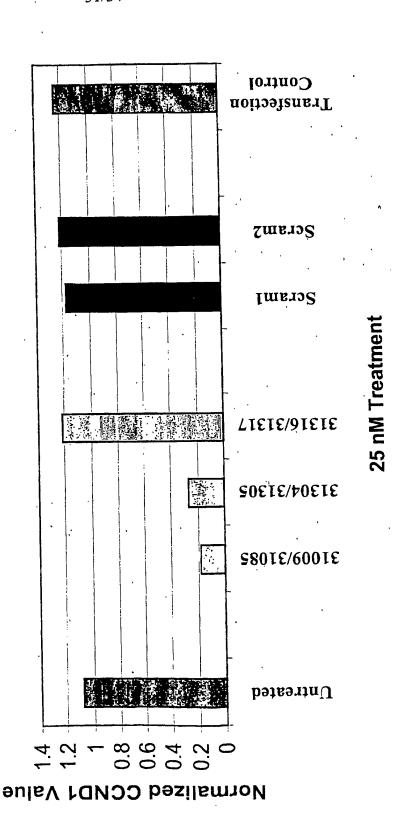


Figure 32: A549 24h PTPN1 mRNA Expression

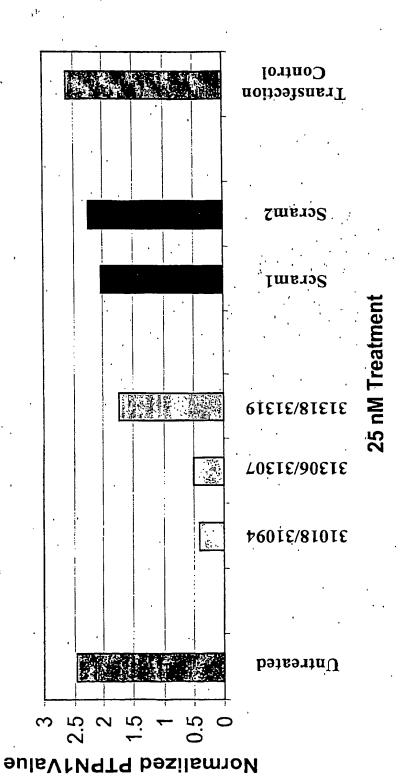


Figure 33: HeLa 24h ERG2 mRNA Expression

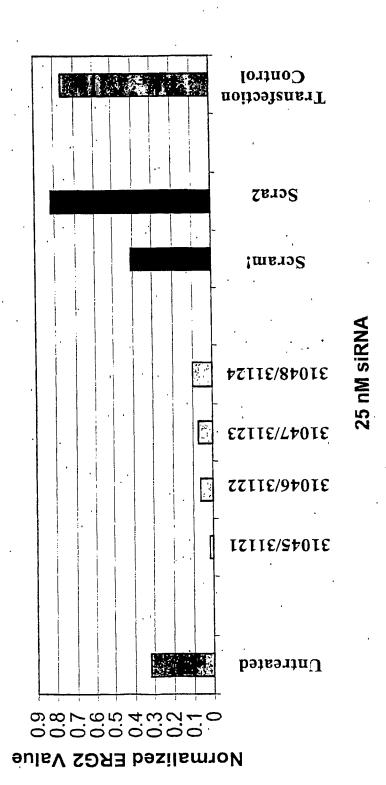
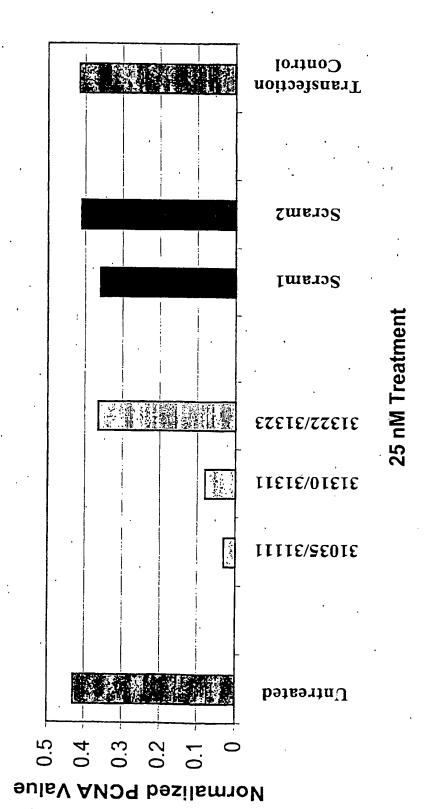


Figure 34: A549 24h PCNA mRNA Expression



		• ,

### (19) World Intellectual Property Organization

International Bureau





(43) International Publication Date 12 September 2003 (12.09.2003)

PCT

### (10) International Publication Number WO 2003/074654 A3

C07H 21/04, (51) International Patent Classification<sup>7</sup>: A61K 48/00, C12N 15/85, 15/86, C12P 19/34, C12Q 1/68

(21) International Application Number:

PCT/US2003/005028

(22) International Filing Date: 20 February 2003 (20.02.2003)

(25) Filing Language:

English

(26) Publication Language:

English

(30) Priority Data:

60/358,58	30 2	0 February	2002 (20	0.02.2002)	US
60/363,12	24	11 March	2002 (1	1.03.2002)	US
60/386,78	82	6 June	2002 (0	6.06.2002)	US
60/406,7	84	29 August	2002 (29	9.08.2002)	US
60/408,3	78 5	September	2002 (0:	5.09.2002)	US
60/409,29	93 9	September	2002 (09	9.09.2002)	US
60/440,13	29	15 January	2003 (1:	5.01.2003)	US

(63) Related by continuation (CON) or continuation-in-part (CIP) to earlier applications:

US	60/358,580 (CON)
Filed on	20 February 2002 (20.02.2002)
US	60/363,124 (CON)
Filed on	11 March 2002 (11.03.2002)
US	60/386,782 (CON)
Filed on	6 June 2002 (06.06.2002)
US	60/406,784 (CON)
Filed on	29 August 2002 (29.08.2002)
US	60/408,378 (CON)
Filed on	5 September 2002 (05.09.2002)
US	60/409,293 (CON)
Filed on	9 September 2002 (09.09.2002)
US	60/440,129 (CON)
Filed on	15 January 2003 (15.01.2003)

(71) Applicant (for all designated States except US): SIRNA THERAPEURICS, INC [US/US]; 2950 Wilderness Place, Boulder, CO 80301 (US).

80301 (US). BEIGELMAN, Leonid [US/US]; 5530 Colt Drive, Longmont, CO 80503 (US). CHOWRIRA, Bharat [US/US]; 576 Manorwood Lane, Louisville, CO 80027 (US). PAVCO, Pamela [US/US]; 705 Barberry Circle, Lafayette, CO 80026 (US). FOSNAUGH, Kathy [US/US]; 1030 Edinboro Drive, Boulder, Colorado 80305 (US). JAMISON, Sharon [US/US]; 4985 Twin Lakes Rd, #89, Boulder, CO 80301 (US). USMAN, Nassim [US/US]; 2129 Night Sky Lane, Lafayette, CO 80026 (US). THOMPSON, James [US/US]; 705 Barberry Circle, Lafayette, CO 80026 (US).

- (74) Agent: TERPSTRA, Anita, J.; McDonnell Boehnen Hulbert & Berghoff, 300 South Wacker Drive, Suite 3200, Chicago, IL 60606 (US).
- (81) Designated States (national): AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CO, CR, CU, CZ, DE, DK, DM, DZ, EC, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, OM, PH, PL, PT, RO, RU, SC, SD, SE, SG, SK, SL, TJ, TM, TN, TR, TT, TZ, UA, UG, US, UZ, VC, VN, YU, ZA, ZM, ZW.
- (84) Designated States (regional): ARIPO patent (GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZM, ZW), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, BG, CH, CY, CZ, DE, DK, EE, ES, FI, FR, GB, GR, HU, IE, IT, LU, MC, NL, PT, SE, SI, SK, TR), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD, TG).

### Declaration under Rule 4.17:

of inventorship (Rule 4.17(iv)) for US only

### **Published:**

with international search report

(88) Date of publication of the international search report: 5 February 2004

For two-letter codes and other abbreviations, refer to the "Guidance Notes on Codes and Abbreviations" appearing at the beginning of each regular issue of the PCT Gazette.

- (72) Inventors; and
- (75) Inventors/Applicants (for US only): MCSWIGGEN, James [US/US]; 4866 Franklin Drive, Boulder, CO

(54) Title: RNA INTERFERENCE MEDIATED INHIBITION OF GENE EXPRESSION USING SHORT INTERFERING NU-CLEIC ACID (SINA)

(57) Abstract: The present invention concerns methods and reagents useful in modulating gene expression in a variety of applications, including use in therapeutic, diagnostic, target validation, and genomic discovery applications. Specifically, the invention relates to small nucleic acid molecules, such as short interfering nucleic acid (siNA), short interfering RNA (siRNA), double-stranded RNA (dsRNA), micro-RNA (miRNA), and short hairpin RNA (shRNA) molecules capable of mediating RNA interference (RNAi) against target nucleic acid sequences. The small nucleic acid molecules are useful in the treatment of any disease or condition that responds to modulation of gene expression or activity in a cell, tissue, or organism.



# INTERNATIONAL SEARCH REPORT

International application No.

PCT/US03/05028

	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1					
TIS CT	IPC(7) : C07H 21/04; A61K 48/00; C12N 15/85, 15/86; C12P 19/34; C12Q 01/68 US CL : 435/6, 91.1, 375; 536/24.5; 514/44					
According to	International Patent Classification (IPC) or to both nat	tional classification and IPC				
	OS SEARCHED umentation searched (classification system followed b	v classification symbols)				
U.S.: 43	5/6, 91.1, 375; 536/24.5; 514/44					
Documentatio	on searched other than minimum documentation to the	extent that such documents are included	d in the fields searched			
Electronic dates	ta base consulted during the international search (name e caplus, lifesci, embase, USPATFULL, biosis	e of data base and, where practicable, s	earch terms used)			
C. DOCU	JMENTS CONSIDERED TO BE RELEVANT					
Category *	Citation of document, with indication, where app	propriate, of the relevant passages	Relevant to claim No.			
X 	WO 00/44914 A1 (MEDICAL COLLEGE OF GEOI INC.) 03 August 2000 (03.08.00), see entire docum	RGIA RESEARCH INSTITUTE, nent.	1-11, 27-29, 32-35, 51-54			
Y			12-26, 30, 31, 36-50			
Y	US 5, 814,620 A (ROBINSON ET AL.) 29 Septemb document.	per 1998 (29.09.98), see entire	1-36, 51-54			
Y, P	WO 02/22636 A1 (ISIS PHARMACEUTICALS, IN	C.) 21 March 2002, see entire	1-36, 42, 51-54			
document Y,P FUTAMI ET AL. Induction of apoptosis in HeLa cells with siRNA expression vector targeted against bcl-2. Nucleic Acids Research Supplement. January 2002, No. 2., pages 251-252, see entire document.			1-35, 41 and 51-54			
X,P  Y,P	TUSCHL ET AL. Small Interfering RNAs: A Revolution and Gene Therapy. Molecular Interverses 158-167, see entire document.	lutionary Tool for the Analysis of rentions. June 2002, Vol. 295, No. 3,	1-11, 27-29, 31-35, 41, 51-54 			
Furthe	r documents are listed in the continuation of Box C.	See patent family annex.	1			
* 5	Special categories of cited documents:  It defining the general state of the art which is not considered to be	"T" later document published after the ir date and not in conflict with the app principle or theory underlying the in	lication but cited to understand the			
of particular relevance  "E" earlier application or patent published on or after the international filing date		"X" document of particular relevance; the considered novel or cannot be considered novel or tannot be considered novel to taken alone	particular relevance; the claimed invention cannot be ovel or cannot be considered to involve an inventive step			
"L" document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)						
"P" documer	at referring to an oral disclosure, use, exhibition or other means of published prior to the international filing date but later than the	"&" document member of the same pate				
Date of the actual completion of the international search  Date of mailing of the international search report						
	3 (15.06.2003)					
	Name and maning address of the form					
Co	ommissioner for Patents	Karen A. Lacourciere				
Al	O. Box 1450 exandria, Virginia 22313-1450	Telephone No. (703) 308-0196	101			
	Jo. (703)305-3230 SA/210 (second sheet) (July 1998)		-U			

PCT/US03/05028

### INTERNATIONAL SEARCH REPORT

Hischemical and biophysical Research Communications. June 2002, Vol. 295, pages 744-748.  LIN ET AL. A Novel mRNA-cDNA Interference Phenomenon for Silencing bcl-2 Expression in Human LNCaP Cells. Biochemical and Biophysical Research Communications. January 2001, Vol. 281, pages 639-644, see entire document.  FIRE ET AL. Potent and specific genetic interference by double-stranded RNA in Caenorhabditis elegans. Nature. 19 February 1998, Vol 391, pages 806-811.  LISENSHIR ET AL. Duplexes of 21-nucleotide RNAs mediate RNA interference in cultured mammalian cells. Nature. 24 May 2001, Vol. 411, pages 494-498, see entire document.  TUSCHL ET AL. Targeted mRNA degradation by double-stranded RNA in vitro. Genes and Development. 15 December 1999, Vol. 13, No. 24, pages 3191-3197, see entire document.  TUSCHL ET AL. Functional anatomy of siRNAs for mediating efficient RNAi in Drosophila melanogaster embryo lysate. The EMBO Journal. October 2001, Vol. 20, No. 23, pages 6877-6888, see entire document.  LUS 6,506,559 B1 (FIRE ET AL.) 14 January 2003 (14.01.03), see entire document.  CA 2359180 A1 (KREUTZER ET AL.) 08 March 2000 (03.08.00), see entire document.  WO 01/36646 A1 (CANCER RESEARCH CAMPAIGN TECHNOLOGY LIMITED) 25 May 2001 (25.05.01), see entire document.  WO 94/01550 A1 (HYBRIDON, INC.) 20 January 1994 (20.01.94), see entire document.	levant to claim No
X LIN ET AL. A Novel mRNA-cDNA Interference Phenomenon for Silencing bcl-2 Expression in Human LNCaP Cells. Biochemical and Biophysical Research Communications. January 2001, Vol. 281, pages 639-644, see entire document.  FIRE ET AL. Potent and specific genetic interference by double-stranded RNA in Caenorhabditis elegans. Nature. 19 February 1998, Vol 391, pages 806-811.  ELBASHIR ET AL. Duplexes of 21-nucleotide RNAs mediate RNA interference in cultured mammalian cells. Nature. 24 May 2001, Vol. 411, pages 494-498, see entire document.  TUSCHL ET AL. Targeted mRNA degradation by double-stranded RNA in vitro. Genes and Development. 15 December 1999, Vol. 13, No. 24, pages 3191-3197, see entire document.  Y,P ELBASHIR ET AL. Functional anatomy of siRNAs for mediating efficient RNAi in Drosophila melanogaster embryo lysate. The EMBO Journal. October 2001, Vol. 20, No. 23, pages 6877-6888, see entire document.  A,P US 6,506,559 B1 (FIRE ET AL.) 14 January 2003 (14.01.03), see entire document.  X CA 2359180 A1 (KREUTZER ET AL.) 08 March 2000 (03.08.00), see entire document.  Y,P WO 01/36646 A1 (CANCER RESEARCH CAMPAIGN TECHNOLOGY LIMITED) 25 May 2001 (25.05.01), see entire document.  X WO 94/01550 A1 (HYBRIDON, INC.) 20 January 1994 (20.01.94), see entire document.  Y WO 99/32619 A1 (THE CARNEGIE INSTITUTE OF WASHINGTON) 01 July 1999 (01.07.99), see entire document.  Y WO 99/4029 A1 (AG-GENE AUSTRALIA LIMITED) 30 September 1999 (30.09.99),	1, 27-29, 31-3: 49, 51-54
Expression in Human LNCaP Cells. Biochemical and Biophysical Research Communications. January 2001, Vol. 281, pages 639-644, see entire document.  Z FIRE ET AL. Potent and specific genetic interference by double-stranded RNA in Caenorhabditis elegans. Nature. 19 February 1998, Vol 391, pages 806-811.  Z ELBASHIR ET AL. Duplexes of 21-nucleotide RNAs mediate RNA interference in cultured mammalian cells. Nature. 24 May 2001, Vol. 411, pages 494-498, see entire document.  Z TUSCHL ET AL. Targeted mRNA degradation by double-stranded RNA in vitro. Genes and Development. 15 December 1999, Vol. 13, No. 24, pages 3191-3197, see entire document.  Y,P ELBASHIR ET AL. Functional anatomy of siRNAs for mediating efficient RNAi in Drosophila melanogaster embryo lysate. The EMBO Journal. October 2001, Vol. 20, No. 23, pages 6877-6888, see entire document.  A,P US 6,506,559 B1 (FIRE ET AL.) 14 January 2003 (14.01.03), see entire document.  X CA 2359180 A1 (KREUTZER ET AL.) 08 March 2000 (03.08.00), see entire document.  Y WO 01/36646 A1 (CANCER RESEARCH CAMPAIGN TECHNOLOGY LIMITED) 25 May 2001 (25.05.01), see entire document.  X WO 94/01550 A1 (HYBRIDON, INC.) 20 January 1994 (20.01.94), see entire document.  Y WO 99/32619 A1 (THE CARNEGIE INSTITUTE OF WASHINGTON) 01 July 1999 (01.07.99), see entire document.  Y WO 99/49029 A1 (AG-GENE AUSTRALIA LIMITED) 30 September 1999 (30.09.99),	28, 30, 36-48,
I2-26,  X FIRE ET AL. Potent and specific genetic interference by double-stranded RNA in Caenorhabditis elegans. Nature. 19 February 1998, Vol 391, pages 806-811.  X ELBASHIR ET AL. Duplexes of 21-nucleotide RNAs mediate RNA interference in cultured mammalian cells. Nature. 24 May 2001, Vol. 411, pages 494-498, see entire document.  X TUSCHL ET AL. Targeted mRNA degradation by double-stranded RNA in vitro. Genes and Development. 15 December 1999, Vol. 13, No. 24, pages 3191-3197, see entire document.  Y,P ELBASHIR ET AL. Functional anatomy of siRNAs for mediating efficient RNAi in Drosophila melanogaster embryo lysate. The EMBO Journal. October 2001, Vol. 20, No. 23, pages 6877-6888, see entire document.  A,P US 6,506,559 B1 (FIRE ET AL.) 14 January 2003 (14.01.03), see entire document.  CA 2359180 A1 (KREUTZER ET AL.) 08 March 2000 (03.08.00), see entire document.  Y,P WO 01/36646 A1 (CANCER RESEARCH CAMPAIGN TECHNOLOGY LIMITED) 25 May 2001 (25.05.01), see entire document.  WO 94/01550 A1 (HYBRIDON, INC.) 20 January 1994 (20.01.94), see entire document.  Y WO 99/32619 A1 (THE CARNEGIE INSTITUTE OF WASHINGTON) 01 July 1999 (01.07.99), see entire document.  WO 99/49029 A1 (AG-GENE AUSTRALIA LIMITED) 30 September 1999 (30.09.99),	1, 27-29, 31-3: 41, 51-54
Caenorhabditis elegans. Nature. 19 February 1998, Vol 391, pages 806-811.  ELBASHIR ET AL. Duplexes of 21-nucleotide RNAs mediate RNA interference in cultured mammalian cells. Nature. 24 May 2001, Vol. 411, pages 494-498, see entire document.  TUSCHL ET AL. Targeted mRNA degradation by double-stranded RNA in vitro. Genes and Development. 15 December 1999, Vol. 13, No. 24, pages 3191-3197, see entire document.  Y.P. ELBASHIR ET AL. Functional anatomy of siRNAs for mediating efficient RNAi in Drosophila melanogaster embryo lysate. The EMBO Journal. October 2001, Vol. 20, No. 23, pages 6877-6888, see entire document.  X. CA 2359180 A1 (FIRE ET AL.) 14 January 2003 (14.01.03), see entire document.  X. CA 2359180 A1 (KREUTZER ET AL.) 08 March 2000 (03.08.00), see entire document.  Y.P. WO 01/36646 A1 (CANCER RESEARCH CAMPAIGN TECHNOLOGY LIMITED) 25 May 2001 (25.05.01), see entire document.  X. WO 94/01550 A1 (HYBRIDON, INC.) 20 January 1994 (20.01.94), see entire document.  Y. WO 99/32619 A1 (THE CARNEGIE INSTITUTE OF WASHINGTON) 01 July 1999 (01.07.99), see entire document.  Y. WO 99/49029 A1 (AG-GENE AUSTRALIA LIMITED) 30 September 1999 (30.09.99),	26, 30, 36-40, 4 50
X ELBASHIR ET AL. Duplexes of 21-nucleotide RNAs mediate RNA interference in cultured mammalian cells. Nature. 24 May 2001, Vol. 411, pages 494-498, see entire document.  X TUSCHL ET AL. Targeted mRNA degradation by double-stranded RNA in vitro. Genes and Development. 15 December 1999, Vol. 13, No. 24, pages 3191-3197, see entire document.  Y, P ELBASHIR ET AL. Functional anatomy of siRNAs for mediating efficient RNAi in Drosophila melanogaster embryo lysate. The EMBO Journal. October 2001, Vol. 20, No. 23, pages 6877-6888, see entire document.  A,P US 6,506,559 B1 (FIRE ET AL.) 14 January 2003 (14.01.03), see entire document.  X CA 2359180 A1 (KREUTZER ET AL.) 08 March 2000 (03.08.00), see entire document.  Y,P WO 01/36646 A1 (CANCER RESEARCH CAMPAIGN TECHNOLOGY LIMITED) 25 May 2001 (25.05.01), see entire document.  X WO 94/01550 A1 (HYBRIDON, INC.) 20 January 1994 (20.01.94), see entire document.  Y WO 99/32619 A1 (THE CARNEGIE INSTITUTE OF WASHINGTON) 01 July 1999 (01.07.99), see entire document.  Y WO 99/49029 A1 (AG-GENE AUSTRALIA LIMITED) 30 September 1999 (30.09.99),	-11, 27-30, 32
cultured mammalian cells. Nature. 24 May 2001, Vol. 411, pages 494-498, see entire document.  X TUSCHL ET AL. Targeted mRNA degradation by double-stranded RNA in vitro. Genes and Development. 15 December 1999, Vol. 13, No. 24, pages 3191-3197, see entire document.  Y, P ELBASHIR ET AL. Functional anatomy of siRNAs for mediating efficient RNAi in Drosophila melanogaster embryo lysate. The EMBO Journal. October 2001, Vol. 20, No. 23, pages 6877-6888, see entire document.  A,P US 6,506,559 B1 (FIRE ET AL.) 14 January 2003 (14.01.03), see entire document.  X CA 2359180 A1 (KREUTZER ET AL.) 08 March 2000 (03.08.00), see entire document.  Y,P WO 01/36646 A1 (CANCER RESEARCH CAMPAIGN TECHNOLOGY LIMITED) 25 May 2001 (25.05.01), see entire document.  X WO 94/01550 A1 (HYBRIDON, INC.) 20 January 1994 (20.01.94), see entire document.  Y WO 99/32619 A1 (THE CARNEGIE INSTITUTE OF WASHINGTON) 01 July 1999 (01.07.99), see entire document.  WO 99/49029 A1 (AG-GENE AUSTRALIA LIMITED) 30 September 1999 (30.09.99),	2-26, 31, 33-54
Y document.  X TUSCHL ET AL. Targeted mRNA degradation by double-stranded RNA in vitro. Genes and Development. 15 December 1999, Vol. 13, No. 24, pages 3191-3197, see entire document.  Y, P ELBASHIR ET AL. Functional anatomy of siRNAs for mediating efficient RNAi in Drosophila melanogaster embryo lysate. The EMBO Journal. October 2001, Vol. 20, No. 23, pages 6877-6888, see entire document.  A,P US 6,506,559 B1 (FIRE ET AL.) 14 January 2003 (14.01.03), see entire document.  X CA 2359180 A1 (KREUTZER ET AL.) 08 March 2000 (03.08.00), see entire document.  Y,P WO 01/36646 A1 (CANCER RESEARCH CAMPAIGN TECHNOLOGY LIMITED) 25 May 2001 (25.05.01), see entire document.  X WO 94/01550 A1 (HYBRIDON, INC.) 20 January 1994 (20.01.94), see entire document.  Y WO 99/32619 A1 (THE CARNEGIE INSTITUTE OF WASHINGTON) 01 July 1999 (01.07.99), see entire document.  Y WO 99/49029 A1 (AG-GENE AUSTRALIA LIMITED) 30 September 1999 (30.09.99),	1-12, 27-35
and Development. 15 December 1999, Vol. 13, No. 24, pages 3191-3197, see entire document.  Y,P ELBASHIR ET AL. Functional anatomy of siRNAs for mediating efficient RNAi in Drosophila melanogaster embryo lysate. The EMBO Journal. October 2001, Vol. 20, No. 23, pages 6877-6888, see entire document.  A,P US 6,506,559 B1 (FIRE ET AL.) 14 January 2003 (14.01.03), see entire document.  X CA 2359180 A1 (KREUTZER ET AL.) 08 March 2000 (03.08.00), see entire document.  Y,P WO 01/36646 A1 (CANCER RESEARCH CAMPAIGN TECHNOLOGY LIMITED) 25 May 2001 (25.05.01), see entire document.  X WO 94/01550 A1 (HYBRIDON, INC.) 20 January 1994 (20.01.94), see entire document.  Y WO 99/32619 A1 (THE CARNEGIE INSTITUTE OF WASHINGTON) 01 July 1999 (01.07.99), see entire document.  Y WO 99/49029 A1 (AG-GENE AUSTRALIA LIMITED) 30 September 1999 (30.09.99),	13-26, 36-54
Y document.  Y,P ELBASHIR ET AL. Functional anatomy of siRNAs for mediating efficient RNAi in Drosophila melanogaster embryo lysate. The EMBO Journal. October 2001, Vol. 20, No. 23, pages 6877-6888, see entire document.  A,P US 6,506,559 B1 (FIRE ET AL.) 14 January 2003 (14.01.03), see entire document.  X CA 2359180 A1 (KREUTZER ET AL.) 08 March 2000 (03.08.00), see entire document.  Y,P WO 01/36646 A1 (CANCER RESEARCH CAMPAIGN TECHNOLOGY LIMITED) 25 May 2001 (25.05.01), see entire document.  X WO 94/01550 A1 (HYBRIDON, INC.) 20 January 1994 (20.01.94), see entire document.  Y WO 99/32619 A1 (THE CARNEGIE INSTITUTE OF WASHINGTON) 01 July 1999 (01.07.99), see entire document.  Y WO 99/49029 A1 (AG-GENE AUSTRALIA LIMITED) 30 September 1999 (30.09.99),	1, 23, 24, 27-3
Drosophila melanogaster embryo lysate. The EMBO Journal. October 2001, Vol. 20, No. 23, pages 6877-6888, see entire document.  A,P US 6,506,559 B1 (FIRE ET AL.) 14 January 2003 (14.01.03), see entire document.  X CA 2359180 A1 (KREUTZER ET AL.) 08 March 2000 (03.08.00), see entire document.  Y,P WO 01/36646 A1 (CANCER RESEARCH CAMPAIGN TECHNOLOGY LIMITED) 25 May 2001 (25.05.01), see entire document.  X WO 94/01550 A1 (HYBRIDON, INC.) 20 January 1994 (20.01.94), see entire document.  Y WO 99/32619 A1 (THE CARNEGIE INSTITUTE OF WASHINGTON) 01 July 1999 (01.07.99), see entire document.  Y WO 99/49029 A1 (AG-GENE AUSTRALIA LIMITED) 30 September 1999 (30.09.99),	22, 25, 26, 31-
X       CA 2359180 A1 (KREUTZER ET AL.) 08 March 2000 (03.08.00), see entire document.       1-         Y,P       WO 01/36646 A1 (CANCER RESEARCH CAMPAIGN TECHNOLOGY LIMITED) 25 May 2001 (25.05.01), see entire document.       1-         X       WO 94/01550 A1 (HYBRIDON, INC.) 20 January 1994 (20.01.94), see entire document.       1-         Y       WO 99/32619 A1 (THE CARNEGIE INSTITUTE OF WASHINGTON) 01 July 1999 (01.07.99), see entire document.       1-         Y       WO 99/49029 A1 (AG-GENE AUSTRALIA LIMITED) 30 September 1999 (30.09.99),	1-54
Y, P WO 01/36646 A1 (CANCER RESEARCH CAMPAIGN TECHNOLOGY LIMITED) 25 May 2001 (25.05.01), see entire document.  X WO 94/01550 A1 (HYBRIDON, INC.) 20 January 1994 (20.01.94), see entire document.  Y  WO 99/32619 A1 (THE CARNEGIE INSTITUTE OF WASHINGTON) 01 July 1999 (01.07.99), see entire document.  Y  WO 99/49029 A1 (AG-GENE AUSTRALIA LIMITED) 30 September 1999 (30.09.99),	1-54
Y,P       WO 01/36646 A1 (CANCER RESEARCH CAMPAIGN TECHNOLOGY LIMITED) 25         May 2001 (25.05.01), see entire document.       1-         X       WO 94/01550 A1 (HYBRIDON, INC.) 20 January 1994 (20.01.94), see entire document.       1-         Y       WO 99/32619 A1 (THE CARNEGIE INSTITUTE OF WASHINGTON) 01 July 1999 (01.07.99), see entire document.       1-         Y       WO 99/49029 A1 (AG-GENE AUSTRALIA LIMITED) 30 September 1999 (30.09.99),	1-35, 51-54
May 2001 (25.05.01), see entire document.  X	36-50
Y  X WO 99/32619 A1 (THE CARNEGIE INSTITUTE OF WASHINGTON) 01 July 1999 1- (01.07.99), see entire document.  Y  WO 99/49029 A1 (AG-GENE AUSTRALIA LIMITED) 30 September 1999 (30.09.99),	1-54
Y  X  WO 99/32619 A1 (THE CARNEGIE INSTITUTE OF WASHINGTON) 01 July 1999 1- (01.07.99), see entire document.  Y  WO 99/49029 A1 (AG-GENE AUSTRALIA LIMITED) 30 September 1999 (30.09.99),	1-35, 51-54
Y (01.07.99), see entire document.  Y WO 99/49029 A1 (AG-GENE AUSTRALIA LIMITED) 30 September 1999 (30.09.99),	36-50
Y WO 99/49029 A1 (AG-GENE AUSTRALIA LIMITED) 30 September 1999 (30.09.99),	1-35, 51-54
(10 001 11 (10 001 10 001 00 00 00 00 00 00 00 00 00	36-50
	1-54

Form PCT/ISA/210 (second sheet) (July 1998)



# This Page is Inserted by IFW Indexing and Scanning Operations and is not part of the Official Record

## **BEST AVAILABLE IMAGES**

Defective images within this document are accurate representations of the original documents submitted by the applicant.

Defects in the images include but are not limited to the items checked:

□ BLACK BORDERS
□ IMAGE CUT OFF AT TOP, BOTTOM OR SIDES
□ FADED TEXT OR DRAWING
□ BLURRED OR ILLEGIBLE TEXT OR DRAWING
□ SKEWED/SLANTED IMAGES
□ COLOR OR BLACK AND WHITE PHOTOGRAPHS
□ GRAY SCALE DOCUMENTS
□ LINES OR MARKS ON ORIGINAL DOCUMENT
□ REFERENCE(S) OR EXHIBIT(S) SUBMITTED ARE POOR QUALITY

# IMAGES ARE BEST AVAILABLE COPY.

OTHER: \_\_\_\_

As rescanning these documents will not correct the image problems checked, please do not report these problems to the IFW Image Problem Mailbox.